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ALIGNMENTS

Human; lymphotoxin beta; receptor; blocking agent; extracellular; ligand binding; domain; treatment; Th1 cell; immune response; delayed; hypersensitivity; contact; tuberculin; granulomatous; graft versus host; disease; organ rejection; autoimmune; disorder; multiple sclerosis; insulin dependent diabetes; uveitis; cytokine; sympathetic ophthalmia; psoriasis; Listeria; Toxoplasma; infection; Mycobacterium; abnormal; lymphoid organ; development. WO9703687-A1. Homo Extracellular domain of human lymphotoxin beta receptor. 29-0CT-1997 AAW23220; AAW23220 sapi ens. standard; (first protein; entry) 197 AA

06-FEB-19 97.

19-JUL-1996; 96WO-US012010.

21-JUL-19 95; 95US-00505606.

(BIOJ) B IOGEN INC

Browning JL, Benjamin CD, Hochman PS;

WPI; 1997 -132373/12.

Composition to treat a diabetes, auto:immune ons comprising lymphotoxin-beta receptor blocking agent auto:immune diseases, e.g. sclerosis, insulin-dependent etc. used

Example μ Page 55-56; 76pp; English.

The presen blocking a human LT-b Th1 cell m hypersensi granulomat rejection dependent agent, com-beta-R up mediated i agent, comprises the extracellular ligand binding domain of the beta-R up to the transmembrane region. It can be used to treat a mediated immune response which contributes to a delayed type sitivity reaction, preferably contact, tuberculin type or tous hypersensitivity, graft versus host disease, organ or an autoimmune disorder, i.e. multiple sclerosis, insulin diabetes, sympathetic ophthalmia, uveitis and psoriasis. It can can മ

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                                                                                                                                                                                                                                                         Human; genoxin; antiarteriosclerotic; antidiabetic; hypotensive; antilipaemic; anorectic; immunomodulator; cytostatic; anti-HIV; antiinflammatory; cardiant; cerebroprotective; gene therapy; tumour necrosis factor receptor; TNFR; body mass; weight loss; obesity.
                                                                                                 W02003011322-A1
                                                                                                                                                                                                                                          Homo sapiens
  Lucas
                                       02-AUG-2001;
                                                           31-JUL-2002;
                                                                                                                               Domain
                                                                                                                                                                      Domain
                                                                                                                                                                                          Protein
                                                                                                                                                                                                             Peptide
                                                                                                                                                                                                                                                                                                                                                      ABR40220;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       signalling e.g., cancer of the prostate, ovary, gallbladder, breast, brain, liver or colon, or inflammatory disorders, such as rheumatoid arthritis, asthma or ulcerative colitis. The present sequence represents a human TNFR2 related protein/LTRbeta amino acid sequence, which is given in comparison with human TNFR2PV in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      and immunosuppressive activities, and can be used TNFR2PV cDNA or protein sequences can be used for for treating a disease or condition associated wit signalling e.g., cancer of the prostate, ovary, gabrain, liver or colon, or inflammatory disorders,
                                                                                                                                                                                                                                                                                                                Human
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disorders
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N-PSDB; F
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                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Screening for an agonist or antagonist of Genoxin activity, useful for preventing or treating metabolic disorders, comprises contacting Genoxin polypeptide with a test compound and determining binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            activity
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ABZ99578.
                                                                                                                                                                                                                                                                                                                                                                          or antagonist of Genoxin. The method is useful in metabolic, particularly, in discovering compounds that modulate Genoxin or that reduce or increase body mass and maintain weight loss, reventing or treating obesity-related diseases or disorders such lipidaemia, atherosclerosis, heart disease, stroke, insulint diabetes or hypertension, or for preventing or treating such as associated with excessive weight loss, such as cachexia, cancer weight loss, acquired immunodeficiency syndrome (AIDS)-related oss, chronic inflammatory disease-related weight loss, or The present sequence represents the human genoxin of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  st of Genoxin activity. The agonists/antagonists of the invention iarteriosclerotic, antidiabetic, hypotensive, antilipaemic, c, immunomodulator, cytostatic, anti-HIV, antiinflammatory, and cerebroprotective activity. The polypeptides of the n may have a use in gene therapy, and act as tumour necrosis sceptor (TNFR) agonists. The method is used to screen for an prantagonist of Genoxin. The method is useful in metabolic particularly, in discovering commounds that it is not as the screen for an activity, in discovering commounds the screen for an activity, in discovering commounds the screen for an activity, in discovering commounds.
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TTCKNPLEPLPPEMSGT 224
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                TTCKNPLEPLPPEMSGT 197
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larity 100.0%;
Conservative (
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Pred. No. 2.3e-78;
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                                                                                                                                                                                                                                                                                                                                                                                                                     Human; cytostatic; DAPK3-Agonist; DAPK3-Antagonist; cancer; TNF-receptor associated factor 5 interacting protein; tumour necrosis factor associated factor 5 interacting protein; TRAF5 interacting protein.
                                                                                                                                                                                     WO2003031
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Best Local
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09-OCT-2001;

09-OCT-2001;

12-OCT-2001;

15-OCT-2001;

17-OCT-2001;

22-OCT-2001;

24-OCT-2001;

29-OCT-2001;

29-OCT-2001;

29-OCT-2001;

01-NOV-2001;

25-JUN-2002;

01-OCT-2002;
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Edinger SR
Mezes PS,
Taupier RJ
                                                                                                                                                                                                                                                                                             The invention describes an isolated polypeptide comprising any of 33 9(1273 amino acid sequences (I) given in the specification or its mature form, a sequence that is at least 95 % identical to (I), or a sequence comprising one or more conservative substitutions in the amino acid sequence of (I). The polypeptide is useful for preparing a composition for treating or preventing e.g. cancer. This is the amino acid sequence of a tumour necrosis factor (TNF)-receptor associated factor 5 (TRAF5) interacting protein associated with the identification of novel human proteins and their functions
                                                                                                                                                                                                                                                                                                                                                                                                                New
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N-PSDB; ACA90237.
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ABP96136
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TRJ,
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2001US-032849P.
2001US-0329414P.
2001US-0341058P.
2001US-0343629P.
2001US-0349575P.
2001US-0346357P.
2002US-0391342P.
2002US-0391342P.
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Zhong H,
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Pred. No. 2.3e-78;
Mismatches 0;
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(TRAF5)
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RESULT 7
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Cytostatic

Antiarteriosclerotic;

Anti-HIV;

Antiinflammatory;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; tumour necrosis factor receptor 2 related protein variant; TNFR2PV; cytostatic; immunosuppressive; antiasthmatic; gene therapy; TNF signalling; cancer; inflammatory disorder; rheumatoid arthritis; asthma; ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                   New cDNA, condition
                                                                                                                                                                                                                                                         The present sequence represents human tumour necrosis factor receptor 2 related protein variant (TNFR2PV). TNFR2PV has cytostatic, antiasthmatic and immunosuppressive activities, and can be used in gene therapy. The TNFR2PV cDNA or protein sequences can be used for preparing a composition for treating a disease or condition associated with increased TNF signalling e.g., cancer of the prostate, ovary, gallbladder, breast, brain, liver or colon, or inflammatory disorders, such as rheumatoid arthritis, asthma or ulcerative colitis
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-25
N-PSDB; ABZ7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (INCY-)
                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                     Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PG,
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                      181
                                              148
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                                                          CPPGTEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSD
                                                                                                           YNEHWNYLTICQLCRPCDPVMGLEBIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSD
                                                                                                                                                               OS.
                                             YNEHWNYLTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSD
                                                                                                                                                                                                                                                                                                                                                                                           useful for preparing a composition for treating a disease associated with increased TNF signaling e.g., cancer of the ovary, gallbladder, breast, brain, liver or colon, or arthritis, asthma.
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79717.
                                                                                                                                       99.6%;
ilarity 99.5%;
Conservative 1
                      CKNPLEPLPPEMSGT 197
CKNPLEPLPPEMSGS
                                                                                                                                                                                                                                        9 AA;
                                                                                                                                                                                                                                                                                                                                                                     g lA-F; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TE GENOMICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US023684.
  224
                                                                                                                                                                                          1;
                                                                                                                                                                                        Score 1129; DB 6;
Pred. No. 4.3e-78;
l; Mismatches 0
                                                                                                                                                                                            0
                                                                                                                                                                                                                  Length 399;
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Matches 192
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27-JUL-2001;
02-AUG-2001;
10-AUG-2001;
10-AUG-2001;
10-AUG-2001;
24-AUG-2001;
31-AUG-2001;
07-SEP-2001;
                                                                                                                                                                                                                                                 cardiovascular, metabolic, developmental, endocrine, muscle, gastrointestinal, lipid metabolism or transport disorders, and viral infections. These are also useful in assessing the effects of exogenous compounds on the expression of nucleic acids and amino acid sequences of REMAP, in facilitating drug discovery process, and in investigating the pathogenesis of diseases or medical conditions. Expression and purification were achieved using bacterial or virus-based expression systems. The present sequence represents an REMAP protein of the invention.
                                                                                                                                                                                                                                                                                                                                         The present invention relates to an isolated polypeptide. The polypeptides and polynucleotides are useful in diagnosing, the preventing disorders associated with aberrant expression of as cell proliferative, autoimmune/inflammatory, renal, neuro cardiovascular, metabolic, developmental, endocrine, muscle,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lal PG, Honchell CD, Forsythe IJ, Walia I Barroso I, Yue H, Warren BA, Thangavelu I Lee EA, Baughn MR, Gorvad AE, Duggan BM, Richardson TW, Elliott VS, Zebarjadian Y, Peterson DP, Luo W, Lehr-Mason PM;
                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                diagnosing, treating of REMAP expression, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                       stroke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antiallergic;
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                                                                                                                                                                               97.8%;
Il Similarity 100.0%;
192; Conservative
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PLEPLPPEMSGT
                                                     EAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCKN
                                                                                     NYLTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPGT
                                                                                                                                  SEQ
                                                                                                                                                                                                                             416
                                           EAELKDEVGKGNNHCVPCKAGHFQNTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                treating or p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2001US-0306020P.
2001US-0308179P.
2001US-0309702P.
2001US-0311476P.
2001US-0311551P.
2001US-0311718P.
2001US-0314798P.
2001US-0316639P.
2001US-0317996P.
                                                                                                                                                                                                                             AA;
                                                                                                                                                                                                                                                                                                                                                                                                               ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2002WO-US022833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antidiabetic;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16;
                     197
                                                                                                                                                                                                                                                                                                                                                                                                                                              l membrane-associated proteins (REMAP), useful
c preventing disorders associated with aberrant
cancer, AIDS, atherosclerosis, hypertension or
                                                                                                                                                                                                                                                                                                                                                                                                              115pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INC
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                                                                                                                                                                              Score 1108; D
Pred. No. 1.8
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pathogenesis.
                                           PSARCQPHTRCENQGLVEAAPGTAQSDTTCKN
                                                                                                                                                                                          DB 7
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Tran UK,
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Gietzen
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JX;
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                                                                                                                                                                                                                                                                                                                                                      neurological
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RESULT 8
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ADA49700
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Matches
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23-SEP-1996;
31-MAR-1997;
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The present
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Apo-2LI, i
receptor h
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Apo-2LI ar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    region of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fluoropho
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-DEC-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-NOV-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             polypeptide wni
ignostic assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      isolated biologically active Apo-2 ligand inhibitor polypeptide, or polypeptide which induces or stimulates apoptotic activity, useful
    21
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                                                                                                                                                                                                                                                                                                       170
                                                                                                                                                     75
                                                                                                                                                                                                \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                igand inhibitor (Apo-2LI) or Apo-3, and the polynucleotide ses encoding them. Apo-2LI and Apo-3 are involved in apoptosis. The and Apo-3 polypeptides are useful in diagnostic assays. Apo-2LI and Apo-3 polypeptides are useful in diagnostic assays. Apo-2LI and for generating antibodies, as standards in assays for Apo-3 or in affinity purification techniques, and in competitive-type shores. Agonistic Apo-3 antibodies are useful for stimulating or gapoptosis in cancer cells, and thus have therapeutic utility. Seent sequence represents the extracellular region of a human TNF of human Apo-2LI.
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KGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCK 184
                                                                                                 RPCDPVMGLEBIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPGTEAELKDEVG
                                                                                                                                                                                                  87.1%; imilarity 100.0%; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ENENTECH INC
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                                                                                                                                                                                                                                                                                                                                                                                                        170 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     region of human TNFRrp (hTNFRrp) protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   96US-00625328.
96US-00710802.
97US-00828683.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53pp; English.
                                                                                                                                                                                                                                                                                                  Score 987; DB 6; I
; Pred. No. 1.2e-67;
0; Mismatches 0;
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                                                                                                                                                                           neuroprotective, antiviral, antiinflammatory, anticonvolution, antiparasitic, cardiant, anti-HIV, antiparkinsonian and vasotropic activities, and can be used in gene therapy. The TRID polynucleotides are useful in the treatment of tumours, resistance to polynucleotides are useful in the treatment of tumours, resistance to parasite, bacteria and viruses, restenosis and graft versus host disease. They are also useful for inducing proliferation of T-cells, endothelial cells and to prevent certain autoimmune diseases after stimulation of TRID by and to prevent certain autoimmune diseases after stimulation of TRID by an agonist or TRAIL binding facilitator. The antibodies which bind TRID cassociated with increased or decreased apoptotic cell death. The TRID polynucleotides, proteins, antibodies, agonists and antagonists are useful in the diagnosis, treatment or prevention of: (a) cancer; (b) autoimmune disorders; (c) diseases associated with increased apoptosis; cautoimmune disorders; and (e) viral infection. The present concerned to the present at the present at the present at the present and the present at the present invention.
                                                                                      Query Match
Best Local S
Matches 135
                                                                                                                                                                                                                                                                                                                                                                                                               The present factor (TNF) related aperintracellular domain, also referred intracellular domain, also referred 5 (TNFR-5 or TR5)). TRID has cytostatic, antiviral, antiinflamma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; tumour necrosis factor receptor 5; TRID; TNFR-5; TR5; nootropic TRAIL receptor without intracellular domain; diagnosis; cytostatic; tumour necrosis factor related apoptosis inducing ligand; vasotropic; immunosuppressive; neuroprotective; antiviral; antiinflammatory; anticonvulsant; antiparasitic; cardiant; anti-HIV; antiparkinsonian; gene therapy; restenosis; graft versus host disease; tumour; cancer; apoptotic cell death related disease; autoimmune disorder; cardiovascular disorder; viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB36700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleic acid encoding a TRID polypeptide, also referred to as t necrosis factor receptor 5, useful in the diagnosis, treatment prevention of cancer, autoimmune disorders and viral infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-MAY-2000;
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                                                                                                                                                    Sequence
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 61
                              28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HUMAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ruben SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                     Similarity
                            415
  YNEHWNYLTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fig 2; 285pp;
                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          necrosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     related apoptosis inducing ligand ('comain, also referred to as tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2000WO-US013515
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                                                                                                                                                     AA;
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                                                                                                        68.0%;
70.7%;
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                                                                                         14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         receptor
                                                                                         Score 771; DB 4;
Pred. No. 8.4e-51;
1; Mismatches 40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the human TRID protein inducing ligand (TRAIL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        immunosuppressive,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         protein
                                                                                            40;
                                                                                                                         Length 41
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          necrosis factor receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ
                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (tumour
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                                                                                                                                                                                                   present in comparison
                                                                                                                                                                                                                                                                                                                                                                                                                                                             nootropic
                                                                                             2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       necrosis
r without
                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ropic;
                                                                                                                                                                                                                                                                                                                                                                     disease
        -ELL
        118
                                   87
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В

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YNEHWNYLTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHC--ELL

118

87

POLVPPYRIENOTOWDODKEYYEPMHDVCCSRCPPGEFVFAVCSRSQDTVCKTCPHNS

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RESULT 10
ABP96138
XX
AC ABP96
AC O9-MI
XX
AC Human
KW TNFR2
CC Mus 1
XX
AC Mus 1
AC Mus 1
XX
AC Mus 1
AC Mu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
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            Ş
                                                                Query Match
Best Local
Matches 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TNFR2PV; cyto
TNF signallin
asthma; ulcen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        condition a prostate, o rheumatoid
                                                                                                                                                                                                                      The present invention describes human tumour necrosis factor receptor related protein variant (TNFR2PV). TNFR2PV has cytostatic, antiasthmat: and immunosuppressive activities, and can be used in gene therapy. The TNFR2PV cDNA or protein sequences can be used for preparing a compositifier treating a disease or condition associated with increased TNF signalling e.g., cancer of the prostate, ovary, gallbladder, breast, brain, liver or colon, or inflammatory disorders, such as rheumatoid arthritis, asthma or ulcerative colitis. The present sequence represent a mouse lymphotoxin-beta receptor amino acid sequence, which is given i comparison with human TNFR2PV in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABP96138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABP9613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-JUL-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INCY-)
                                                                                                                                                                            Sequence
                                                                                                                          Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lymphotoxin-beta receptor protein
                                                                    ll Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tumour necrosis factor receptor 2 related protein variant;
7; cytostatic; immunosuppressive; antiasthmatic; gene therapy;
ynalling; cancer; inflammatory disorder; rheumatoid arthritis;
; ulcerative colitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             œ
                  \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INCYTE GENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SDCPPGTEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQ 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SDTICKNPPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SDTTCKNPLEP 189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Waj
SQPQAVPPYASENQTCRDQEKEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ø
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       seful for preparing a composition for treating a disease ssociated with increased TNF signaling e.g., cancer of thywary, gallbladder, breast, brain, liver or colon, or arthritis, asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                56445/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fig 2A-C;
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2002WO-US023684.
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                                                                                                                                                                                AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          protein;
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                                                                     68.0%; Score 771; DB 6; 70.7%; Pred. No. 8.4e-51; ive 14; Mismatches 40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO:20.
                                                                           40;
                                                                                                                                Length 415;
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RESULT 11
AAW94642
ID AAW94
XX
AC AAW94
XX
AC AAW94
XX
AC TUMO
XX
TUMO
KW TUMO
KW Infla
KW Skin
KW Skin
KW Skin
KW Homo
XX
Homo
XX
Homo
XX
PR W098!
XX
PR W098!
XX
PF 29-MJ
XX
PF 29-MJ
XX
PF 29-MJ
XX
PF 1 Grees
XX
PF 29-MJ
XX
CC TNF
CC Corr
CC Corr
CC Corr
CC Gesi
CC TNF
CC Gesi
CC Gesi
CC Gesi
CC Gomg/k
CC Goma
XX
Sequ
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                                                      Query Match
Best Local S
Matches 77
                                                                                                                                                 correspond in primary sequence to a binding loop of a tumour necrosis factor receptor (TNF-R) superfamily member. The compounds are especially designed from a binding loop of TNF-R p55. They are capable of inhibiting TNF binding to its cellular receptors and may be used to inhibit the biological activities of TNF. They may be used in treating TNF-associated conditions such as acute and chronic inflammatory responses, septic shock, cachexia, autoimmunity, graft-versus-host disease, skin allergic reactions, immune complex disease, transplantation rejection and malaria. Administration is, e.g. oral, transdermal, transmucosal, pulmonary, subcutaneous, intravenous or intramuscular. Parenteral dosage is 0.1-5 mg/kg/day. The present sequence represents an extracellular Cys-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo
                                                                                                                                                                                                                                                                                                                                                                         New compounds designed receptor - are capable necrosis factor, e.g.,
                                                                                                                                             mg/kg/day.
domain of
                                                                                                                                                                                                                                                                                        The present invention d correspond in primary s factor receptor (TNF-R)
                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          30-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tumour necrosis factor receptor inflammation; septic shock; car skin allergic reaction; immune transplantation rejection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TNF-R
                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Greene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               extracellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88
                            52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               VINU
                                                      1 40.2% Similarity 100.0 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               77
VCATCAENSYNEHWNYLTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ഗ -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SDTTCKNPLEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DTICKNEPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Murali
                                                                                                                                                                                                                                                                                                                                                Fig 1; 78pp;
                                                                                                                 AA;
                                                                                                                                                                                                                                                                                                                     invention describes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98WO-US010891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-00866545
                                                                                                                                           from the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cys-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            218
                                                                                                                                                                                                                                                                                                                                                                           from a binding loop of a tumour necrosis factor of inhibiting the biological activities of tumour in treating inflammation or autoimmune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Takasaki
                                                                                                                                                                                                                                                                                                                                                English.
                                                                    0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ptor; TNF-R; autoimmune diseases;
  cachexia; graft versus host diseas
une complex disease; malaria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     77
                                                        ; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             domain TNF-R-rp.
                                                                                  S
                                                      Score 456; DB Pred. No. 1.5; Mismatches
                                                                                                                                                                                                                                                                                        peptides and peptide to a binding loop of a nily member. The compo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Σ,
                                                                                                                                             invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TNF-R;
                                                                    DB
1.5
                                                                      5e-
IAPCTSKRKTQCRCQPGMFCAAWALE
                                                                                     N
                                                                    27;
                                                         0
                                                                                  Length
                                                         Indels
                                                                                                                                                                                                                                                                                                                    analogues which
                                                                                 77;
                                                        0,
                                                      Gaps
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52

Similarity

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Gaps

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RESULT 12
AAB69194
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Query Match
Best Local
Matches
                                                                                             osteoclastogenesis and bone resorption. Osteoclastogenesis and bone resorption inhibiting peptide analogues from the present invention have osteopathic, cytostatic, antirheumatic, antiarthritic, antiinflammatory and immunomodulatory activities, and are tumour necrosis factor (TNF)-related activation-induced cytokine (TRANCE) / receptor activator of NF-kappaB ligand (RANK) inhibitors. The method is useful for treating disease, metastatic bone disease, rheumatoid arthritis or periodontal disease, and modulating dendritic cell maturation, T cell proliferation, and/or CD40 receptor systems. The present sequence represents an extracellular Cys-rich domain of a tumour necrosis factor receptor (TNF-R) superfamily member, which is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       osteoclastogenesis; bone loss; bone resorption; osteoclast; antirheumatic; antiarthritic; antiinflammatory; immunomodulatory; tumour necrosis factor-related activation-induced cytokine; TRANCE; receptor activator of NF-kappaB ligand; RANK; osteoporosis; Paget's disease; metastatic bone disease; rheumatoid arthritis; periodontal disease; modulating dendritic cell maturation; T cell proliferation; CD40 receptor system.
                                                                                                                                                                                                                                                                                                                                         Use
                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                  Aoki
                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYPE-)
(AOKI/)
(HORN/)
                                                                                  present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB69194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAB69194
                                                       Sequence
                                                                                                                                                                                                                                                                                                Disclosure; Fig 1; 81pp; English.
                                                                                                                                                                                                                                                                                                                           Use of peptides and peptide analogs which are TRANCE/RANK inhibitors, for inhibiting osteoclastogenesis and bone resorption.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-JUL-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-APR-20
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                                                                                                                                                                                                                                                                                                                                                                                                                              HORNE
BARON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UNIV PENNSYLVANIA
                                                                                    invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                          OKI K.
                                                                                                                                                                                                                                                                     int invention describes a method for inhibiting
                                                                                                                                                                                                                                                                                                                                                                                                 Horne WC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CTHCELLSDCPPGTEAE
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                                                                                                                                                                                                                                                                                                                                                                       -182866/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
                                                          AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     extracellular Cys-rich domain TNF-R-rp
40.2%; Score 456; DB 4; 1.larity 100.0%; Pred. No. 1.5e-27; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2000WO-US020510.
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C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein;
                                                                                                                                                                                                                                                                                                                                                                                                  Baron R,
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encompasses polypeptides 90% identical and polynucleotides 95% identical compasses polypeptides 90% identical and polynucleotides 95% identical compositions. The invention additionally relates to recombinant vectors and host cells comprising human ovarian antigen composing or preventing polypeptides in diagnosing, treating, prognosing or preventing various ovary and/or breast-related disorders. Such conditions include ovarian cancer and breast-related disorders (e.g., infertility, disorders of pregnancy, anovulation, collections (e.g., chlammatory conditions (e.g., mastities, endocrine disorders, infections (e.g., chlammydia, HIV, toxoplasmosis, and toxic shock syndrome), inflammatory conditions (e.g., mastitis, oophoritis and vaginitis), immune disorders (e.g., congenital and acquired immunodeficiencies, autoimmune oophoritis, systemic lupus erythematosus), blood-related disorders (e.g., anaemia), cardiovascular disorders, condulate ovarian antigen expression or activity. The polynucleotides may also be used in screening for compounds which modulate ovarian antigen expression or activity. The polynucleotides may condenical and in forensic analysis, and the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ovarian cancer; breast cancer; tumour; reproductive system disorder; infertility; pregnancy disorder; anovulation; polycystic ovary syndrome; pCOS; ovarian cyst; dysmenorrhoea; endocrine disorder; infection; inflammatory condition; immune disorder; blood disorder; infection; cardiovascular disorder; respiratory disorder; neurological disorder; gastrointestinal disorder; urinary system disorder; drug screening; gene therapy; chromosome mapping; forensic analysis; antibody preparation; cytostatic; immunomodulatory; neuroprotective; antiinflammatory; gynaecological; reproductive; chromosome 12p13.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isolated nucleic acid molecules encoding novel ovarian polypeptides, useful in the prevention, treatment and diagnosis of cancer (e.g. ov cancer), immune disorders, cardiovascular disorders and neurological
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N-PSDB; ABQ55003.
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(ABQ54131-ABQ56305), and also
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RESULT 14
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Matches 71
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TNFR/fc f
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07-OCT-1994
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AAQ45224 is cDNA from clone 1 of library WI-26 VA4 of human fibroblast cell line WI-26 VA4. The mature full-length TNFRI is a glycoprotein having a mol. wt. of about 75-80 kDa.The cloning of the cDNA for TNFRI was described in Smith et al., Science 248:1019,1990. Clone 1 is contained in expression vector pCAV/NOT-TNFR (ATCC 68088). A recombinant chimeric antibody may be produced having TNFR sequences substituted for the variable domains of either or both of the immunoglobulin molecule heavy and light chains and having unmodified constant region domains. A specific example of a TNFR/Fc fusion protein is given in AAQ45225/R51003 The rhu TNFR:Fc fusion gene was created by ligating the following fragments into a cloning vector: 1) an 867 bp Asp718-Pvu2 fragment from pCAV/NOT-TNFR (ATCC 68088) contg. the cDNA encoding the truncated TNFR. 2) a 700 bp Sty1-Spe1 fragment from plasmid pIXY498 coding for 232 AAs of the coding point of the cDNA encoding the truncated TNFR.
                                                                                                                                                                                                                                                                                                        WPI; 1994-1161/2
N-PSDB; AAQ45225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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sease diagnosis, drug targeting and phenotyping. The
presents a human ovarian antigen of the invention. Not
a for this patent did not form part of the printed
on, but was obtained in electronic format directly from
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                                                                                                                                                                                                                      Page 32-34; 47pp;
                                                                                                                                                                                                                                                   mediated inflammatory diseases with TNF antagonist of TNF receptor, opt. as fusion protein with human in Fc region, esp. for treating arthritis.
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The invention soluble domai biologically
                                                                                                                 New concatameric protein diagnosing and treating dits glycosylated form, suarthritis and cachexia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antiinflammatory; antibacterial; immunosuppressive; antirheum; antiarthritic; immunomodulator; concatameric protein; soluble dimeric protein; inflammation; septicaemia; cytotoxicity; rheumatoid arthritis; cachexia; inflammation; human.
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JC4302
I37552
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A42086
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 17-Mar-2000
C;Accession: I54182
R;Baens, M.; Chaffanet, M.; Cassiman, J.J.; Van den Berghe, H.; Marynen, P.
Genomics 16, 214-218, 1993
A;Title: Construction and evaluation of a hncDNA library of human 12p transcrib.
A;Reference number: I54182; MUID:93252381; PMID:8486360
A;Accession: I54182
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: mRNA
A;Residues: 1-435 <RES>
A;Cross-references: GB:L04270; NID:g339761; PIDN:AAA36757.1; PID:g339762
C;Genetics:
A;Gene: GDB:LTBR
A;Cross-references: GDB:1230195; OMIM:600979
A;Map position: 12p13.3-12p13.1
C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat homology.
RESULT 2
A35356
tumor necrosis factor receptor N; Alternate names: 75K tumor ne C; Species: Homo sapiens (man) C; Date: 10-Sep-1999 #sequence receptor C; Accession: A35356; A36475; A4 R; Smith, C.A.; Davis, T.; Ander Science 248, 1019-1023, 1990 A; Title: A receptor for tumor n
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                  p-1999 #sequence_revision 10-Sep-1999 #text_change 27-Oct-2003
A35356; A36475; A48416; A36007; A23666; B35010; I38094
; Davis, T.; Anderson, D.; Solam, L.; Beckmann, M.P.; Jerzy, R
1019-1023, 1990
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A;Residues; 1-461 c8MY-
A;Cocas-references; 08,103215; NID;gla9185; PIDN;AAA5929.1; PID;gl89186
A;Cocas-references; 08,10;S.A87, 1831-835, 1990
A;Title: A second tumor necrosis factor receptor gene product can shed a;
A;Reference number: A56475 MUD;9104991; PMI:3172993
A;Reference number: A56475 MUD;9104991; PMI:317299
A;Cocas-references; 08:45594; 08:398549; NID;9339757; PIDN;AAA35755.1; P;
R;Dembic, Z.; 1.oetscher, H.; Gubler, U.; Ran, Y.C.; Lahm, H.W.; Gentr., R.;
A;Title: Two human TWP receptors have similar extracellular, but distinct
A;Title: Two human TWP receptors have similar extracellular, but distinct
A;Title: Two human TWP receptors have similar extracellular, but distinct
A;Title: Two human TWP receptors have similar extracellular, but distinct
A;Title: Two human TWP receptors have similar extracellular, but distinct
A;Reference number: A64016; MUD;91370690; PMID;936649
A;Arcession: A48416
A;Reference number: A8416; MUD;91370690; PMID;96649
A;Arcession: A48016
A;Arcession: A46007
A;Arcession: A4600
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A; Status: pre
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nm, H.W.; Gentz,
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tumor necrossing mus mus C; Species: Mus mus C; Date: 30-Jun-199 C; Accession: B3867 R; Lewis, M.; Tartana Nacad.
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A; Residues: 1-474 <LEW>
A; Cross-references: GB:Mi
R; Goodwin, R.G.; Anderson
Mol. Cell. Biol. 11, 3021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Accession type: |
A; Molecule type: |
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C; Superfamil:
C; Keywords:
F; 1-22/Domai:
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A;Cross-references: GB:M60469; NID:g199827; PIDN:AAA39752.1; PID:g199828
R;Kissonerghis, M.; Fellowes, R.; Feldmann, M.; Chernajovsky, Y.
submitted to the EMBL Data Library, May 1995
A;Description: Characterization of the promoter region of the murine p75-A;Reference number: S54816
A;Reference number: S54816
A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-22 <KIS>
PMDI:g809043; PIDN:CAA60618.1; PID:g80904
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F;280-461/Domain
F;171,193/Bindin
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A; Accession: B38
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66-203/Domain
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ar cloning and expression of the type 1 and ter: A40254; MUID:91246168; PMID:1645445
JTICQLCR---PCDPVMGLEEIAPCTSKRKTQCRCQPGMFCA--AWALECTHCELLSDC
                                      KPEPGYECQISQ-EYYDRKAQMCCAKCPPGQYVKHFCNKTSDTVCADCEASMYTQVWN
                                                                    ASE-NOTCRDQEKEYYEPOHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWN
                                                                                                                                                                                                                                  NGF receptor repeat homology <NG1>
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                                                                                                                                  ilarity
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larity 35.0%;
Conservative 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mRNA
                                                                                                                     Conservative
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intracellular #status predicted <INT>
site: carbohydrate (Asn) (covalent) #
                                                                                                                                                                                                                 receptor repeat homology <NG2>
                                                                                                                                                                                             receptor
                                                                                                                 26.0%; Score 295; DB 2; 34.0%; Pred. No. 3.5e-15; ive 24; Mismatches 84;
                                                                                                                                                                                           repeat
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Pred. No. 6.1e-16;
Nismatches 78; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                192
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                                                                                                                                                         Length 474;
                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type
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, C.I.; Cope
                                                                                                                       20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            murine
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121

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C;Date: U.C;Accession: I48854
C;Accession: I48854
R;Powell, E.E.; Wicker, L.S.; Pt.
Mamm. Genome 5, 726-727, 1994
A-Title: Allelic variation of tl
                                                      Α; C:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Residues:
A; Cross-refe
C; Superfamil
F; 151-188/Do
                                                                                                  A; Accession: S04
A; Molecule type:
                                                                                                                A; Reference number: A; Accession: S04460
                                                                                                                                                      EMBO
                                                                                                                                                                                                                                     A6077
                              A; Title: Biochemical characteristics and partial amino A; Reference number: A60771; MUID:89093941; PMID:2463309
                                                                                          A;Residues:
                                                                                                                                          A; Title:
                                                                                                                                                            C;Species: Homo sapiens (man)
C;Date: 03-Jun-1993 #sequence_revision
C;Accession: S04460; A60771
R;Stamenkovic, I.; Clark, E.A.; Seed, I
                                                                                                                                                                                                        B-cell activation protein CD40 precursor N;Alternate names: B-cell surface antigen
                     A; Accession:
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C;Spe
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           ;Molecule
                                                  Cross-references: EMBL:X60592; Braesch-Andersen, S.; Paulie, Immunol. 142, 562-567, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Species: Mus musculus (hor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-references: EMBL:X76401; NID:g433830; Superfamily: tumor necrosis factor receptor 151-188/Domain: NGF receptor repeat homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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1801
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                                                                                                                         8, 1403-1410, 1989

B-lymphocyte act: A B-lymphocyte act: S04460;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match
type: ]: 21-50
                                                                                                                                                                                                                                                                                                                                 131
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                                                                                                                                                                                                                                                                                                                                                                              OFRICLSCSSSCSTDQV
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                                                                                                                                                                                                                                                                                                                                                                                                                                               PYASE-NOTCRDQEKEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPGTEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDT
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                                                                                                    mRNA
        protein
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<BRA>
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                                                                                                                                                                                                                                                                                  APESPTLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor
                                                                                                                          activation molecule 460; MUID:89356608;
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MUID:95178848; PMI
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                                                                            NID: 92985
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                                                                Koho,
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                                                                                                                          PMID:247534
                                                                                                                                                                                                                      human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; DB 2; L
8.2e-15;
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PMID:7873884
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                                                                            PIDN: CAA43045.1;
                                                                Nika,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      type 2
y <NGF>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PIDN:CAA53981.1;
type 2 (TNFR2);
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                                        sequence
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NGF receptor
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em, P.;
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                                                             Perlmann,
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                                                                                                                                   receptor
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A;Gene: GDB:CD40
A;Cross-references: GDB:215268; OMIM:109535
A;Cross-references: GDB:215268; OMIM:109535
A;Map position: 20q12-20q13.2
C;Superfamily: CD27 antigen; NGF receptor repeat homology C;Keywords: B-cell; glycoprotein; phosphoprotein; surface F;1-20/Domain: signal sequence #status predicted <SIG>F;21-277/Product: B-cell activation protein CD40 #status F;21-193/Domain: extracellular #status predicted <EXT>F;194-215/Domain: transmembrane #status predicted <TMM>F;153,180/Binding site: carbohydrate (Asn) (covalent) #st
                                                                                                                                                       C; Comment: For C; Superfamily: C; Keywords: alt F; 105-144/Domai
                                                                                                                                                                                                                                                                                                                       A; Accession: A46515
A; Status: preliminary; not
A; Molecule type: nucleic ac
                                                                                                                                                                                                                 C; Comment:
                                                                                                                                                                                                                                                                                        A;Residues: 1-287,'
                                                                                                                                                                                                                                                                                                                                                                                 R;Grimaldi, J.C.; Torres, R.; Kozak, C.A.; Chang, R.; Claz
J. Immunol. 149, 3921-3926, 1992
A;Title: Genomic structure and chromosomal mapping of the
A;Reference number: A46515; MUID:93094586; PMID:1281194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Residues: 1-305 < A; Cross-references:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J. Immunol. 148, 620-626, 1992
A;Title: Differential increase of an alternatively polyadenylated mRNA
A;Reference number: A46476; MUID:92105763; PMID:1370315
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A46476
B cell-
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C;Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               A;Note:
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Best Loc
Matches
                                                                                                  Local
                                                                                                                  Match
                                                                                                                                                                                                                                                                                                                                                                                                                                    -references: GB:M83312; NID:g1553058
sequence extracted from NCBI backbone (NCBIN:75206, NCBIP:75207)
this translation is not annotated in GenBank entry MUSCD40A, rel
ldi, J.C.; Torres, R.; Kozak, C.A.; Chang, R.; Clark, E.A.; Howar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
                                                                                                                                                                                                                                                   sequence
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18-Jun-1993 #sequence_revision
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TCSD--KQYLHDGQ--CCDLCQPGSRLTSHCTALEKTQCHPCDSGEFSAQWNREIRCHQH
                                                                                                                                                    an alternative splice form, see PIR:A46515.

an alternative splice form, see PIR:A46476.

CD27 antigen; NGF receptor repeat homology ternative splicing; transmembrane protein in: NGF receptor repeat homology <NGF>
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-cell; glycoprotein; phosphoprotein; surface antigen; transmembrane
: signal sequence #status predicted <SIG>
uct: B-cell_activation protein CD40 #status experimental <MAT>
                                                                                               imilarity
                                                                                                                                                                                                                                                                                                            287,'LV' <GRI>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ted surface molecule CD40,
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                                     CRDQEKEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWNYLTICQLC
                                                                                                                                                                                                                                                                 source: BALB/c, liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Clark, E.A.
, 620-626, 1992
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                                                                                                                                                                                                                                                  extracted from NCBI backbone (NCBIP:120357)
                                                                              Conservative
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intracellular #status predicted <CYT>
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                                                                                          21.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24.5%;
                                                                                                                                                                                                                                                                                                                                               compared with conceptual
                                                                                                                                                                                                                                                                                     NID: g1553058; PIDN: AAB08705.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lymphoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20;
                                                                        Score 243.5; DB 2
Pred. No. 1.9e-11;
2; Mismatches 81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 278; DB 2; Pred. No. 4.4e-14;
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                                                                                                               DB 2;
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                                                                                                                305;
                                                                                                                                                                                                                                                                                     PID:g1553059; GB:M94126;
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Graph receptor-6 - chicken
C; Species: Gallus gallus (chicken)
C; Date: 09-Nov-2001 #sequence_revision 09-Nov
C; Accession: JC7705
R; Bridgham, J.T.; Bobe, J.; Goetz, F.W.; John
Biochem. Biophys. Res. Commun. 284, 1109-1115
A; Title: Conservation of death receptor-6 in
A; Reference number: JC7705
A; Accession: JC7705
A; MOlecule type: mRNA
A; Residues: 1-651 <BRI>
A; Cross-references: GB:AF349908
C; Comment: This receptor, a member of the tum
tresia, activates a cell death and/or surviva
C; Genetics:
A; Gene: dr-6
C; Superfamily: tumor necrosis factor receptor
C; Keywords: ovary
F; 1-21/Domain: signal sequence #status predic
F; 332-350/Domain: extracellular cysteine-rich,
F; 332-350/Domain: death domain #status predic
F; 410-475/Domain: conserved cytoplasmic #status
Fredic
R;Mallett, S.; Fossum, S.; Barclay, A.N.

EMBO J. 9, 1063-1068, 1990

A;Title: Characterization of the MRC OX40 antigen of activated A;Reference number: S12783; MUID:90214614; PMID:2157591

A;Recession: S12783

A;Molecule type: mRNA

A;Residues: 1-271 <MAL>
A;Cross-references: EMBL:X17037; NID:g57830; PIDN:CAA34897.1;
C;Superfamily: CD27 antigen; NGF receptor repeat homology
C;Keywords: growth factor receptor; transmembrane protein
F;1-19/Domain: signal sequence #status predicted <MAT>
F;20-271/Product: OX40 antigen #status predicted <TMM>
F;211-235/Domain: transmembrane #status predicted <TMM>
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                                                                                                                                                                                            OX40 antigen precursor - rat
N;Alternate names: nerve growth factor receptor
C;Species: Rattus norvegicus (Norway rat)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-19
C;Accession: S12783; S08036
R;Mallett, S.; Fossum, S.; Barclay, A.N.
EMBO J. 9, 1063-1068, 1990
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extracellular cysteine-rich,
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ptor-6 in avian and pi
308433; PMID:11414698
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No. 6.5e-10;
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N; Alternate:
N; Contains:
C; Species: H
C; Date: 30-J
C; Accession:
R; Fuchs, P.;

30-Jun-

A38208; Strehl,

Homo

sis factor receptor 1 precursor [validated] - human names: P55 tumor necrosis factor receptor; TNF receptor type tumor necrosis factor alpha inhibitor; tumor necrosis factor fomo sapiens (man)
Jun-1992 #sequence_revision 30-Jun-1992 #text change 08-Dec-20

1 binding

prote

A60231;

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#sequence revision 30-Jun-1992 #text_change 08-Dec-2000 A34899; A34900; A36555; C36555; A38281; S12057; JT0758; S.; Dworzak, M.; Himmler, A.; Ambros, P.F.

tumor GQHUT1

necrosis

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gene ox40 protein N;Alternate names: C;Species: Mus mus C;Date: 02-Jul-199 C;Accession: I4870 R;Calderhead, D.M. J. Immunol. 151, 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Accession: I48700
A;Status: translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                               A; Cross-reference
C; Genetics:
                                                                                                                                                                                                                                                                                                                                        A;Status: (A;Molecule
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151, 5261-5271, 1955
10ning of mouse Ox40: 748700; MUID
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                                                                                                                                                                                                       Similarity 31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            KDTYPSGHK-CCRECQPGHGMVSRCDHTRDTVCHPCEPGFYNEAVNYDT-CKQCTQCNHR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SGSELKONCTPTEDTVCQCR-
                                                                  VPCKAGHFQNTSSP--SARCQPHTRCENQGLVEAAPGTAQSDTTCKN
                                                                                                                      MGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPGTEAELKDEVGKGNNHC
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                                                                                                                                                   KEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWNYLTICQLCRPCDPV
                                                                                               SGSELKQNCTPTQDTVCRCR-
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                                                                                                                                                                                                                                                          86/1; 122/1; 144/2; 210/1; D27 antigen; NGF receptor r
                                                                                                                                                                                                                                                                                                                ,'G',16-272 <RE2>
es: EMBL:X85214; NID:g732818; PIDN:CAA59476.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96 #sequence revision 02-Jul-1996 #text
00; I48334; S34377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sculus (house )
                                          CPPGHF--
                                                                                                                                                                                                                                                                                                                                             DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                  <RES>
                                                                                                                                                                                                                                                                                                                                                                               : EMBL:Z21674; NID:g312827; PIDN:CAA79772.1; PID:g312828 Copeland, N.G.; Gilbert, D.J.; Jenkins, N.A.; Barclay, 5, 926-930, 1995 cture and chromosomal localization of the mouse homologue 148334; MUID:95255413; PMID:7737295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mouse
                                          --SPGNNQACKPWTNCTLSGKQTRHPASDSLDAVCED
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                                                                                                                                                                                                                   18.9%; Score 214.5; DB 2 31.7%; Pred. No. 2.7e-09;
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                                                                                                                                                                                                                                     Length 272;
                                                                                                                                                                                                             Indels
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                                                                                                   PGTQPR-QDSGYKLGVDC
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A; Note: L...
A; Note: L...
R; Nophar, Y.; Nophar, Y.;
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A; Residues: 1-13
R; Seckinger, P.;
Eur. J. Immunol.
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A; Residues: 1-455 <HI
A; Cross-references: (
A; Accession: C36555
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A; Cross-refe
R; Loetscher,
Cell 61, 351
                                                                                                                                                                                                                                                                                                                           A;Title: Cloning and partial A;Reference number: JT0758; A;Accession: JT0758
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A; Residues: 30-38; 41-53, 'X', 55-79, 'XX', 82-94,
A; Residues: 10-38; 41-53, 'X', 55-79, 'XX', 82-94,
A; Note: the purified protein, called tumor ne
R; Gray, P.W.; Barrett, K.; Chantry, D.; Turne
Proc. Natl. Acad. Sci. U.S.A. 87, 7380-7384,
A; Title: Cloning of human tumor necrosis fact
                      A; Molecule type: protein
A; Residues: 41-43, 'X', 45-53, 'X', 55-57
R; Gatanaga, T.; Hwang, C.; Kohr, W.; (
Proc. Natl. Acad. Sci. U.S.A. 87, 878
                                                                                                                                                                           A; Title: Tumor necre
A; Reference number:
                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Note: parts of soluble R;Kemper, O.; Wallach, D Gene 134, 209-216, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Residues: 1-455 < A;Cross-references:
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A; Residues: 1-45
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Cell
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A;Cross-references: GB:M37764
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A; Residues: 1-455
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Cell 61, 351-359, 1990
A;Title: Molecular cloning and A;Tit
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A; Molecule type: DNA
A; Molecule type: DNA
- noeidues: 1-455 < FUC>
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NA Cell Biol.
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                                                                                                                                                      Accession:
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, T.J.; Lewis, M.; Koller,
                                                                                                                                                                   ger, P.; Vey, E.; Turcatti, G.; Willimmunol. 20, 1167-1174, 1990
Tumor necrosis factor inhibitor: pnce number: A60231; MUID:90292116;
   ya, T.; Hwang, C.;
:1. Acad. Sci. U.S
Purification and
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Structure
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e number: S12057;
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of the human
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   Kohr, W.; Cappuccini,
3.A. 87, 8781-8784, 199
characterization of an
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MUID:94085779; PM
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Kroenke, M.
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Jahm, H.W.;
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A;Experimental source: cancer patient serum R;Olsson, I.; Lantz, M.; Nilsson, E.; Peetr Eur. J. Haematol. 42, 270-275, 1989
A;Title: Isolation and characterization of A;Reference number: A60594; MUID:89171156;
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ct: TNF binding protein 1 (tumor necrosis
NGF receptor repeat homology <NG1>
n: NGF receptor repeat homology <NG2>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nces: GDB:125913; OMIM:191190
1: 12p13.2-12p13.2
3; 65/1; 108/1; 158/1; 184/2;
                                                                                                                                                       LTICQLCRPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWA---LECTHCELLSDCPPG
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-53,'X',55-144,'X',146-150,'X',152-186,'X',188-201
                               TTCKNPL
                                                                                                                                                                                                                                                                                                                  inding site:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein is one of
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NGF receptor repeat homology <NG4 >
transmembrane #status predicted <MEM>
intracellular #status predicted <INT>
ing site: carbohydrate (Asn) (covalent
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216
                               187
                                                             -KONTVCT-CHAGFFLRENECVSCSNCKKSLECTKLCLPQIENVKGTEDSG
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                                                                                                                                                                                                                                                   19;
                                                                                                                                                                                                                                                    Score 210.5; DB Pred. No. 8e-09; 9; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            two known
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Peetre,
                                                                                                                                                                                                                                                                                                                                                                                                                               (tumor necrosis
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PMID:2924890
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11

F;110-151/Domain:

NGF

receptor repeat homology <NG3>

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A; Recession: D36858
A; Accession: D36858
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-349 < BLI>
A; Residues: 1-349 < BLI>
A; Cross-references: GB: X69198; NID: g456758; PIDN: CAA49137.1; PII
A; Cross-references: GB: X69198; NID: g456758; PIDN: CAA49137.1; PII
A; Cross-references: GB: X69198; NID: g456758; PIDN: CAA49137.1; PII
A; Cross-references: Strain India-1967, ssp. major, isolate Ir
R; Kolykhalov, A.A.; Blinov, V.M.; Gytorov, V.V.; Pozdnyakov, S.C
submitted to the EMBL Data Library, April 1992
A; Reference number: S46868
A; Reference number: S46868
A; Reference number: S46868
A; A; Cross-references: EMBL: X67117; NID: g516428; PIDN: CAA47540.1; l
A; Cross-references: EMBL: X67117; NID: g516428; PIDN: CAA47540.1; l
A; Experimental source: strain India-1967, isolate Ind3
R; Shchelkunov, S.N.; Blinov, V.M.; Sandakhchiev, L.S.
FEBS Lett. 319, 80-83, 1993
A; Reference number: S32385; MUID: 93202281; PMID: 8384129
A; Accession: S32385
A; Accession: S32385
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A; Status: preliminary; translated for A; Molecule type: DNA
A; Residues: 1-348 < MAS>
A; Cross-references: EMBL:L22579; NII
A; Experimental source: strain Bangla C; Superfamily: TNF-alpha receptor-II
A; Accession: S32385
A; Molecule type: DNA
A; Residues: 31-168 <SHC>
A; Cross-references: EMBL:X
A; Experimental source: str
C; Genetics:
A; Gene: G4R
C; Superfamily: TNF-alpha r
F; 32-66/Domain: NGF recept
F; 68-109/Domain: NGF recept
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N;Alternate names: B28R prote:
C;Species: variola virus
C;Date: 30-Sep-1993 #sequence
C;Accession: D36858; S46888;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Blinov, V.M. submitted to GenBank,
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source: strain Bangladesh 1975
TNF-alpha receptor-II; NGF rec
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B28R protein
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ce: strain ]
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ptor repeat h
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S32385; S3
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    homology <
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ed. No. 7.1e-09;
Mismatches 80
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C;Species: va
C;Date: 24-No
C;Accession:
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G2R pro
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A;Description: Analysis of the complete coding sequence of DNA, Reference number: A72150
A;Accession: D72175
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C;Superi
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C; Accession: JC4
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A;Cross-references: GB:Y16780; NID:g5830555;
A;Experimental source: strain Garcia-1966
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                                                                      A; ritle: Cloning of the cDNA encoding the porcine p55 tumor A;Reference number: JC4302; MUID:96011645; PMID:7590278 A;Accession: JC4302
                                                                                                        R;Suter, B.; Pau
Gene 163, 263-26
A;Title: Cloning
                                                                                                                                                                                                                                                                                     Ş
                                                                                                                                                                                                       JC4302
                                                                                                                                                                                                                                                            밁
                                                                                                                                                                                        tumor
                 A;Accession: PC4093
                                       A;Residues: 1-46
A;Cross-referenc
                                                     A; Molecule type:
A; Residues: 1-46
A;Residues:
          A; Molecule
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3, 263-266,
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Pauli, U.
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<6, 1995
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 : protein <SU2>
                                                                                                                                                                                                                                                                                       ELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQP 160
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                                                    1 <SUT>
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                                          NID:g1141752;
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                                              PIDN:AAC48499.1;
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                                              PID:g1141753
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A;Experimental source: kidney cell line 15

C;Genetics:
A;Gene: tnfr

C;Superfamily: tumor necrosis factor receptor type 1; NGF receptor repeat PC;Keywords: glycoprotein; kidney; receptor; transmembrane protein; tumor F;1-29/Domain: signal sequence #status predicted <SIG>
F;30-461/Product: tumor necrosis factor receptor p55 #status predicted <MAPF;44-194/Domain: extracellular cysteine rich #status predicted <MAPF;44-82/Domain: NGF receptor repeat homology <NGI>
F;84-126/Domain: NGF receptor repeat homology <NGF>
F;361-447/Domain: transmembrane #status predicted <SIT>
F;361-447/Domain: signal transduction #status predicted <SIT>
F;54,145,151/Binding site: carbohydrate (Asn) (covalent) #status predicted
Search completed: Job time : 42 sec
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I37552

OX40 homolog - human
C; Species: Homo sapiens (man)
C; Date: 29-May-1998 #sequence_revision 29-May-1998 #text_cha
C; Date: 29-May-1998 #sequence_revision 29-May-1998 #text_cha
C; Accession: I37552
R; Latza, U.; Durkop, H.; Schnittger, S.; Ringeling, J.; Eite
Eur. J. Immunol. 24, 677-683, 1994
A; Title: The human OX40 homolog: cDNA structure, expression
A; Reference number: I37552; MUID:94170844; PMID:7510240
A; Accession: I37552
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A;Molecule type: mRNA
A;Residues: 1-277 <RES>
A;Cross-references: EMBL:X75962;
C;Superfamily: CD27 antigen; NGF
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Similarity 28.3%;
19; Conservative :
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                        2004,
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                                                                                                                                                                                                                                                                                                                                                                                                                           Score 201; DB
Pred. No. 2.8e-
21; Mismatches
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med. No. 1.8e-08;
Mismatches 77;
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.08;
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FUNCTION. MEDLINE=94225209; PubMed=8171323; Crowe P.D., VanArsdale T.L., Walter B.N., Ware C.F., Hession C., Crowe P.D., VanArsdale T.L., Din W.S., Goodwin R.G., Smith C.A.; Ehrenfels B., Browning J.L., Din W.S., Goodwin R.G., Smith C.A.; "A lymphotoxin-beta-specific receptor."; Science 264:707-710(1994). [4] CHARACTERIZATION.	SEQUENCE FROM N.A. TISSUB=Lung; MEDLINE=22388257; PubMed=12477932; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Richards S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J., Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).	SEQUENCE FROM N.A. SEQUENCE FROM N.A. TISSUE=Liver; MEDLINE=93252381; PubMed=8486360; Baens M., Chaffanet M., Cassiman J.J., den Berghe H., Marynen P.; "Construction and evaluation of a hncDNA library of human 12p transcribed sequences derived from a somatic cell hybrid."; Genomics 16:214-218(1993). [2]	P36941; 01-JUN-1994 (Rel. 29, Created) 01-JUN-1994 (Rel. 29, Last sequence update) 10-OCT-2003 (Rel. 42, Last annotation update) Tumor necrosis factor receptor superfamily member 3 precursor (Lymphotoxin-beta receptor) (Tumor necrosis factor receptor 2 related protein) (Tumor necrosis factor C receptor). LTBR OR TNFRSF3 OR TNFCR. Homo sapiens (Human). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. NCBI_TaxID=9606;], [

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A Mizushima S.-I., Fujita M., Ishida T., Azuma S., Kato K., Hirai M.,

Tumor necrosis factor receptor-associated factor 5 (TRAF5).";

Gene 207:135-140(1998).

-!- FUNCTION: Receptor for the heterotrimeric lymphotoxin containing

-!- FUNCTION: Receptor for the heterotrimeric lymphotoxin containing

-!- FUNCTION: Receptor for the heterotrimeric lymphotoxin containing

-!- FUNCTION: Receptor TNFS14/LIGHT. Promotes apoptosis via TRAF3

C -!- SUBUNIT: Self-associates. Associates with TRAF3, TRAF4 and TRAF5.

-!- SUBCELLULAR LOCATION: Type I membrane protein.

C -!- SIMILARITY: Contains 4 TNFR-Cys repeats.
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MIM; 600979; -.
GO; GO:0007165; P:signal transduct.
GO; GO:0007165; P:signal transduct.
InterPro; IPR008063; Fas_receptor.
InterPro; IPR001368; TNFR_c6.
Pfam; PF00020; TNFR_c6; 4.
PRINTS; PR01680; FASRECEPTOR.
PRINTS; PR01680; TNFR; 4.
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Wu M.-Y., Wang P.-Y., Han S.-H., Hsieh S.-L.;
"The cytoplasmic domain of the lymphotoxin-beta death in HeLa cells.";
J. Biol. Chem. 274:11868-11873(1999).
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"TRAF5, an activator of NF-kappaB
the lymphotoxin-beta receptor.";
J. Biol. Chem. 271:14661-14664(19)
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MEDLINE=98289299; PubMed=9626059;

Krajewska M., Krajewski S., Zapata J.M., Var
Gascoyne R.D., Berern K., McFadden D., Shaba
Reynolds A., Clevenger C.V., Reed J.C.;

"TRAF-4 expression in epithelial progenitor
adult, fetal, and tumor tissues.";

Am. J. Pathol. 152:1549-1561(1998).
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Tumor necrosis factor receptor superfamily member
(Lymphotoxin-beta receptor).
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MEDLINE=7., Tashiro K., Nazarea M.,
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           rap and chromosomal (1995).
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Nakano H., Oshima H., Chung W.,
Yagita H., Okumura K.;
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- FUNCTION: Receptor for the heterotrimeric lymphotoxin cont
LTA and LTB, and for TNFS14/LIGHT. Promotes apoptosis via
and TRAF5. May play a role in the development of lymphoid
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SUBUNIT: Self-associates (By similarity). Associates with TRAF3 and TRAF4 (By similarity)
SUBCELLULAR LOCATION: Type I membrane protein.
SIMILARITY: Contains 4 TNFR-Cys repeats.
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TNFR-CYS 4.
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N-LINKED (GLCNAC...
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Pred. No. 7.1e-5
4; Mismatches
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MEDLINE=22388257; PubMed=12477932;

A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

A Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
                                                                                                                                                                                                                                                                                                                                                      SEQUENCE I
ARG-301.
Rieder M.;
Rieder M.;
Rieder M.;
Schackwitz
Schackwitz
Submitted
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MEDLINE=223
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Lepaslier
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Smith C.
Dower S.
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Mammalia; E
NCBI_TaxID=
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[3]
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MEDLINE=91045991; PubMed=2172983;
Kohno T., Brewer M.T., Baker S.L., Schwartz P.E., King M.W.,
Hale K.K., Squires C.H., Thompson R.C., Vannice J.L.;
"A second tumor necrosis factor receptor gene product can shed naturally occurring tumor necrosis factor inhibitor.";
Proc. Natl. Acad. Sci. U.S.A. 87:8331-8335(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=90260639; PubMed=2160731;
Smith C.A., Davis T., Anderson D., So
Dower S.K., Cosman D., Goodwin R.G.;
"A receptor for tumor necrosis facto:
cellular and viral proteins.";
Science 248:1019-1023(1990).
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3; Q16042; Q9UIH1;
3-1991 (Rel. 17, Created)
3-1991 (Rel. 41, Last sequence update)
8-2003 (Rel. 41, Last sequence update)
R-2004 (Rel. 43, Last annotation update)
necrosis factor receptor superfamily member 1B precursor (Tumor sis factor receptor 2) (p80) (TNF-R2) (p75) (CD120b) (Etanercept)
sis factor necrosis factor binding protein 2 (TBPII)].
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FROM N.A.

96299745; PubMed=8661109;

C.P., White P.S., Maris J.M., Sulman E.P.,

L., Stallard B.J., Goeddel D.V., Desauvage

G.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90260639; Pu
A., Davis T.
                                                                                                                                                                                                                                                                                                                                                     J., Livingston R.J., Daniels M.R., Chung M.-W.,
K.E., Nguyen C.P., Nguyen D.A., Poel C.L., Robertson P.D.,
z W.S., Sherwood J.K., Witrak L.A., Nickerson D.A.;
l (MAR-2003) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                      FROM N.A., AND VARIANTS ARG-196; LYS-232; PRO-269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mapping and genomic structure of the human TNFR2 35:94-100(1996).
                                                                                                                                                                                                                                                                                                                        FROM N.A.
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[ [8]

SEQUENCE OF 154-183 FROM N.A., AND VARIANTS

MEDLINE=21069356; PubMed=11197692;

Tsuchiya N., Komata T., Matsushita M., Oha

T "New single nucleotide polymorphisms in th

TINFR2: association with systemic lupus ery

Genes Immun. 1:501-503(2000).

[9]

RP

MEDLINE=90110215; PubMed=2153136;

Engelmann H., Novick D., Wallach D.;

RT

"Two tumor necrosis factor-binding proteince. Evidence for immunological cross-j

urine. Evidence for immunological cross-j

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Part is a sequence in immunological cross-j

crossis factor receptors.";
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Pennica D., Lam V.T., Mize N
Lipari M.T., Goeddel D.V.;
"Biochemical properties of t
Characterization of ligand b
phosphorylation.";
J. Biol. Chem. 267:21172-211
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SEQUENCE OF 37-461
MEDLINE=91370690;
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"Complementary DNA and demonstration of Proc. Natl. Acad. St
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SEQUENCE OF 116-46:
MEDLINE=90349572; |
Heller R.A., Song |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIAN LOET SCHIBEYER - LOETSCHER H., Schibeyer - Brockhaus M.;
Brockhaus M.;
"Purification and partial amino acid sequence "Purification and partial amino acid sequence distinct tumor necrosis factor receptors from distinct tumor necrosis factor receptors from chem. 265:20131-20138(1990).
                                                                                                                                                              Park Y.C., Bu "Structural k human TRAF2."
                                                                                                                                                                                                                                  TRAF2.
MEDLINE=9
                                                                                                                                                                                                                                                                                                                        INTERACTION WITH TRAF2.

MEDLINE=94349371; PubMed=8069916;

Rothe M., Wong S.C., Henzel W.J., Goeddel D.V.;

"A novel family of putative signal transducers a cytoplasmic domain of the 75 kDa tumor necrosis Cell 78:681-692(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J. Biol. Chem. 26 [11] CHARACTERIZATION.
VARIANTS ARG-196 AND LYS-232.
MEDLINE=21603988; PubMed=11762942;
Morita C., Horiuchi T., Tsukamoto H.,
Arinobu Y., Otsuka T., Sawabe T., Hara
"Association of tumor necrosis factor
                                                                                                                                                                                                                                                                              X-RAY
                                                                                                                                       Nature
                                                                                                                                                                                                                                                                                                    [13]
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C.,_Burkit1
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72; PubMed=21669
ong K., Onasch M
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; PubMed=1966549;
cher H., Gubler U
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tt V., Villa A.R., Tong
s for self-association;
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PubMed=2173696;
laeger E.J., Lahm
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T., Mize N.K.,
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-!- FUNCTION; R
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-!- PTM: A soluble form (tumor necrosis factor binding protein 2) is produced from the membrane form by proteolytic processing.
-!- PHARMACEUTICAL: Available under the name Enbrel (Immunex and Wyeth-Ayerst). Used to treat moderate to servere rheumatoid arthritis (RA). Enbrel consist of the extracellular ligand-binding portion of TNFR2 linked to an Immuglobulin Fc chain. It binds to TNF-alpha and blocks its interactions with receptors.
-!- SIMILARITY: Contains 4 TNFR-Cys repeats.
-!- DATABASE: NAME=PROW; NOTE=CD guide CD120b entry; WWW="http://www.ncbi.nlm.nih.gov/prow/cd/cd120b.htm".
-!- DATABASE: NAME=Enbrel; NOTE=Clinical information on Enbrel; WWW="http://www.enbrel.com/".
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leum. 44:2819-2827(2001).
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lan J.L., Castello R., Moghetti P
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L; BC052977; AAA63262.1; -.
L; M35857; AAA63262.1; -.
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R Kissonerghis M., Fellowes R., Feldmann M., Chernajovsky Y.;

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MEDLINE=91187885; PubMed=1849278;
Lewis M., Tartaglia L.A., Lee A., Benne Wong G.H., Chen E.Y., Goeddel D.V.;

"Cloning and expression of cDNAs for two necrosis factor receptors demonstrate specific.";

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Mol. Cell. Biol.
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01-MAY-1992 (Rel. 22, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Tumor necrosis factor receptor superfamily member necrosis factor receptor 2) (TNF-R2) (p75).
TNFRSF1B OR TNFR2 OR TNFR-2.
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)106 for factor receptor superfamily member of the formula for factor (Decoy receptor 3) (DcR3)
OR DCR3 OR TR6.

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B precursor (M68).

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D=9606;

; Metazoa; Eutheria;

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TISSUE=Fetal lung;
MEDLINE=99087326; PubMed=987
Pitti R.M., Marsters S.A., L
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TISSUE=Prostate;
MEDLINE=99253915; PubMed=10318773;
Yu K.-Y., Kwon B., Ni J., Zhai Y., Ebner R., Kwon B.S.;
"A newly identified member of tumor necrosis factor receptor superfamily (TR6) suppresses LIGHT-mediated apoptosis.";
J. Biol. Chem. 274:13733-13736(1999).
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Godowski P.J., Wood W.I., Gurney A.L., Hill
Goddard A.D., Botstein D., Ashkenazi A.;
"Genomic amplification of a decoy receptor
colon cancer.";
Nature 396:699-703(1998).
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Proc. Natl. Acad. So
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Bai C., Connolly B., Metzker M.L., Hilliard C.A., Liu X., Soderman A., Galloway S.M., Liu Q., Austin C.P., Caskey C. "Overexpression of M68/DcR3 in human gastrointestinal tracindependent of gene amplification and its location in a fo
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EMBL; BC017065; AAH17065.1; -.
EMBL; BC034349; AAH34349.1; -.
HSSP; O14763; 1D0G.
Genew; HGNC:11921; TNFRSF6B.
MIM; 603361; -.
GO; GO:0005625; C:soluble fraction; TAS.GO; GO:0004872; F:receptor activity; TAS.GO; GO:0006916; P:anti-apoptosis; inhibitor act GO; GO:0006916; P:anti-apoptosis; TAS.Ffam; PF00020; TNFR C6; 4.
SMART; SM00208; TNFR C6; 4.
SMART; SM00208; TNFR NGFR 1; 1.
PROSITE; PS00652; TNFR NGFR 2; 2.
ENGRETOR: APOPTOSIS: GTVCOPTOLEIN; Repeat
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                                       (Receptor receptor)
                                           10-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Tumor necrosis factor receptor superfamily mer
(Receptor activator of NF-KB) (Osteoclast diffreeptor) (ODFR).
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TR11_
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                                          RANK.
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TNFR_NGFR_2; 2.
is; Glycoprotein;
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TNFR-CYS 4.
BY SIMILARITY.
N-LINKED (GLCNAC.
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SUPERFAMILY
TNFR-CYS 1.
TNFR-CYS 2.
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Pred. No. 1.5e-17;
3; Mismatches 77
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Mammalia; I
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hi; Muridae; Murinae; Mu
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CTISSUE-Mammary gland;

X MEDLINE=22388257; PubMed=12477932;

X MEDLINE=22388257; PubMed=12477932;

A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

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A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Worley K.C., Hale S., Garcia A.M., Gay L.J., Mullahy S.J.,

A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulky S.W.,

A Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Butterfield Y.S.N., Krzywinski M.I., Schemutz J., Myers R.M.,

B Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schemutz J., Myers R.M.,

B Blakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schemutz J., Myers R.M.,

B Utterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,

Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

T "Generation and initial analysis of more than 15,000 full-length

Thuman and mouse cDNA sequences.";

L Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
EMBL; AF019046; AAB86810.1; -.
EMBL; BC019185; AAH19185.1; -.
HSSP; P25942; 1CDF.

MGD; MGI:1314891; Tnfrsf11a.
GO; GO:0007275; P:development; IM
GO; GO:0007515; P:lymph gland dev
GO; GO:0001503; P:ossification; I
InterPro; IPR001368; TNFR_c6.
Pfam; PF00020; TNFR_c6; 3.
SMART; SM00208; TNFR; 4.
PROSITE; PS00652; TNFR_NGFR_1; 1.
PROSITE; PS50050; TNFR_NGFR_2; 1.
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SEQUENCE FROM N.A.
TISSUE=Fetal liver;
MEDLINE=98032977; PubMed=936
"~~~rson D.M., Maraskovsky E
                                                                                                                                                               EMBL;
EMBL;
HSSP;
MGD; N
GO; G(GO; Q(GO); Q(
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Morinaga T., Higashio K.;
"RANK is the essential signaling receptor for osteocla
differentiation factor in osteoclastogenesis.";
Biochem. Biophys. Res. Commun. 253:395-400(1998).
-!- FUNCTION: Receptor for TNFSF11/RANKL/TRANCE/OPGL;
RANKL-mediated osteoclastogenesis. Involved in the
interactions between T-cells and dendritic cells.
-!- SUBUNIT: Interacts with TRAF1, TRAF2, TRAF3, TRAF1
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Nature 39
[2]
SEQUENCE
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"A homologue of the TNF rece and dendritic-cell function. Nature 390:175-179(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             - SUBUNIT: Interacts with TRAF1, T. similarity).
- SUBCELLULAR LOCATION: Type I mem!
- TISSUE SPECIFICITY: UBIQUITOUS E: TRABECULAR BONE, THYMUS, SMALL I! WEAKLY EXPRESSED IN SPLEEN AND BOUSE SIMILARITY: Contains 4 TNFR-Cys
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PubMed=9367155;
raskovsky E., Billingsley W.L.,
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Stamenkovic I., Clark E.A., Seed B.;
"A B-lymphocyte activation molecule
factor receptor and induced by cytok
EMBO J. 8:1403-1410(1989).
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s P., Matthews L.H., Ashurst J., Stavrides G., Almeida J.P., lJ., Barlow K.F., Bates K.N., Bel O.P., Bird C.P., Blakey S.E., l Burrill W.D., Butler A.P., Ca
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                                                                                                                                                                                                                                                         STANDARD;
                                                                                                                                                                                               22, Last sequence update)
43, Last annotation update)
ctor receptor superfamily member 5 precursor
(B-cell surface antigen CD40) (CDw40) (Bp50)
                                                  PubMed=11780052;
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195
47
61
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234
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POTENTIAL.
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; Pred.
25; Mis
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CYTOPLASMIC (POTENTIAL)
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TNFR-CYS 3.
TNFR-CYS 4.
BY SIMILARITY.
N-LINKED (GLCNAC.
N-LINKED (GLCNAC.
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Catarrhini;
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                                                                                        cytokines in
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F8C1872E99511D8E
            J.P., Burton J., Gilbert J
J.P., Babbage A.K., Bagguley
N., Beard L.M., Beare D.M.,
S.E., Bridgeman A.M., Brown I
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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No. 5e-17;
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    n A.M.,
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A Chapman J.C., Clamp M., Clark G., Clark L.N., Clark S.Y., Clee C.M., A Clegg S., Cobley V.E., Collier R.E., Connor R.E., Corby N.R., A Coulson A., Coville G.J., Deadman R., Dhami P.D., Dunn M., A Ellington A.G., Frankland J.A., Fraser A., French L., Garner P., B Ellington A.G., Frankland J.A., Fraser A., French L., Garner P., A Hammond S., Harley J.L., Heath P.D., Ho S., Holden J.L., Howden P.J., Huckle E., Hunt A.R., Hunt S.E., Jekosch K., Johnson C.M., Johnson D. A., Kimberley A.M., King A., Knights A., Laird G.K., Lawlor S.A., Kay M.P., Kimberley A.M., King A., Knights A., Laird G.K., Lawlor S.A., Martin S.L., McConnachie L.J., McLay K., McMurray A.A., A Milne S.A., Mistry D., Moore M.J.F., Mullikin J.C., Nickerson T., Oliver K., Parker A., Patel R., Pearce T.A.V., Peck A.I., Phillimore B.J.C.T., Prathalingam S.R., Plumb R.W., Ramsay H., A Rice C.M., Ross M.T., Scott C.E., Sehra H.K., Shownkeen R., Sims S., Swann R.M., Sycamore N., Taylor R., Tee L., Thomas D.W., Thorpe A., A Whitehead S.L., Whittaker P., Willey D.L., Williams S.A. Whitehead S.L., Whittaker P., Willey D.L., Williams S.A. Wilming L., Waray P.W., Hubbard T., Durbin R.M., Bentley D.R., Beck S.A.
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Tracey A., L., Why Whitehead S.L., Wray I wilming L., Wray I worders J.;
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SEQUENCE FROM N.A. (ISOFORM II).
SEQUENCE FROM N.A. (ISOFORM II).
MEDLINE=21117110; PubMed=11172023;
Tone M., Tone Y., Fairchild P.J., Wykes M., Waldmann H.;
Tone M., Tone Y., Fairchild P.J., Wykes M., Waldmann H.;
"Regulation of CD40 function by its isoforms generated through "Regulation of CD40 function by its isoforms."
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TISSUE=Ov
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Proc. Natl. Acad. Sci.
[4]
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                                                                                                                                                                                                                                                                                         INTERACTION WITH TRAF3.

MEDLINE=95184010; PubMed=7533327;

Cheng G., Cleary A.M., Ye Z.S., Hong
"Involvement of CRAF1, a relative of
Science 267:1494-1498(1995).
                                                       INTERACTION WITH TRAF1; TRAF2; TMEDLINE=98384149; PubMed=9718306
Pullen S.S., Miller H.G., Everde
                                                                                                                                                proteins b
                    "CD40-tumor necrosis
                                                                                                                                                                                                                                                     INTERACTION WITH TRAF3
   interactions:
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A., Grouse L.H., Derge J.G.,
A., Grouse L.H., Derge J.G.,
Wagner L., Shenmen C.M., Schuler G.D
Juetow K.H., Schaefer C.F., Bhat N.K.,
Juetow K.H., Schaefer C.F., Bhat N.K.,
Juetow K.H., Schaefer C.F., Bhat N.K.,
Juetow K.H., Schaefer C.F.,
Juetow K.H., Schaefer T.,
Jueto
    receptor-associated CD40 signaling throu
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    factor (TRA)
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D., Mullahy S.J.,
Gunaratne P.H.,
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Lovell J.D.,
                          (TRAF)
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n P.J.,
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MEDLINE=98095
Kashiwada M.,
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"Tumor necros
extracellular
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Mizushima S.-I.,
Otsuka M., Yamamo
                                                                                                                                                                                                                                                                                                                                "Downstream
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3D-STRUCTURE MODELING OF 26-186 IN COMPLEX WITH CD40L.
MEDLINE=98266353; PubMed=9605317;
Singh J., Garber E., van Vlijmen H., Karpsusas M., Hsu Y.-M.,
Sheng Z., Naismith J.H., Thomas D.;
"The role of polar interactions in the molecular recognition of CD40L with its receptor CD40.";
Protein Sci. 7:1124-1135/1006\
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MEDLINE=97189482; PubMed=9037712;
Bajorath J., Aruffo A.;
"Construction and analysis of a detailed three-dimensional model of the ligand binding domain of the human B cell receptor CD40.";
Proteins 27:59-70(1997).
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                                                                                                                                                                                                                                             VARIANT HIGM3 I
MEDLINE=2153291
Ferrari S., Gi
Loubser M., Av
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                                                                                                                                                                                                                  Levy Y., Cat
Plebani A.;
                                                                                                                                                                                                                                                                                                                 TRAF3.";
Structure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            X-RAY
                                                                                                                   "Mutations of CD40 gene cause an autosomal recessive immunodeficiency with hyper IgM."; Proc. Natl. Acad. Sci. U.S.A. 98:12614-12619(2001).-!- FUNCTION: Receptor for TNFSF5/CD40L.-!- SUBUNIT: Interacts with TRAF1, TRAF2, TRAF3, TRAF-1-SUBCELLULAR LOCATION: Type I membrane protein (is secreted (isoform II).
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Satterthwait
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Name=II
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TISSUE |
DISEASE
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Event=A.
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ing and characterization of a cDNA encoding the human homolog of
necrosis factor receptor-associated factor 5 (TRAF5).";
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Comment=Additional isoforms seem to exist;
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32985; PubMed=11675497;
32985; PubMed=11675497;
Giliani S., Insalaco A., Al-Ghonaium A., Soresina F
Giliani S., Insalaco M., Badolato R., Ugazio A.G.
Avanzini M.A., Marconi M., Badolato R., Ugazio A.G.
Avanzini M.A., Marconi M., Notarangelo L.D.,
                                                                                                                                                                                                                                                                                                                                                         t A.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                      42386; PubMed=10984535;
lsh K., Leo E., Chiou C.K., Wu H., Reed basis for CD40 signaling mediated by TRA Acad. Sci. U.S.A. 97:10395-10399(2000).
                                                                                                                                                                                                                                                                                                                  0:403-411 (2002) .
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(=P25942-2; Sequence=VSP_006472, VSP_SPECIFICITY: B-cells and in primary: Defects in TNFRSF5 are the cause
                                                                                                             TIVE PRODUCTS:
                                                     =P25942-1; Sequence=Displayed;
                                                                                                                                                                                                                                                                                                                                          Z., Havert M.L., Cabezas E., He J., Kaiser D., Reed J.C., A.C., Cheng G., Ely K.R.; regulator TANK binds to the CD40 recognition site on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -140(1998).
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37:11836-11845(1998).
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3482; PubMed=9037712;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PubMed=9511754;
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Matches
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16-OCT-2001 (Rel. 40, I

10-OCT-2003 (Rel. 42, I

Tumor necrosis factor r

(Receptor activator of

receptor) (ODFR).

TNFRSF11A OR RANK.

Homo sapiens (Human).

Eukaryota; Metazoa; Cho

Mammalia; Eutheria; Pri

NCBI_TaxID=9606;
                                                                     Tometsko
Galibert
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Q9Y6Q6;
16-OCT-2001
                                                                                Anderson D.M., Maraskovsky Tometsko M.E., Roux E.R.,
MEDLINE=99097247;
             FUNCTION
                                                                                                          TISSUE=Dendritic cell; MEDLINE=98032977; PubM
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                                   homologue of the TNF receptor d dendritic-cell function."; ture 390:175-179(1997).
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L; AL035662; CAC17670.1; -.
L; AJ300189; CAC29424.1; -.
L; BC012419; AAH12419.1; -.
; S04460; A60771.
; 1CDF; 01-APR-97.
; 1FLL; 18-OCT-00.
; 1L0A; 08-FEB-00.
; 1CZZ; 26-SEP-01.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cells to undergo isotype switching, one of the find differentiation steps in the humoral immune system to mount an antibody-specific immune response, and germinal center formation.

SIMILARITY: Contains 4 TNFR-Cys repeats.

DATABASE: NAME=PROW; NOTE=CD guide CD40 entry;

WWW="http://www.ncbi.nlm.nih.gov/prow/cd/cd40.htm"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunodeficiency type 3 (HIGM3) [MIM:606843]. HIGM3 autosomal recessive disorder which includes an inabicells to undergo isotype switching, one of the final
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Similarity 35.0%;
2; Conservative
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PubMed=98785
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askovsky E., B
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40, Last sequence update)
42, Last annotation update)
tor receptor superfamily meml
r of NF-KB) (Osteoclast diffo
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Primates;
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0; Mismatches
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Catarrhini
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differentiation facto
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Receptor;
Disease mu
SIGNAL

utation;

Deafness.

TUMOR NECROSIS FACTOR SUPERFAMILY MEMBER 11A

MEMBER 11A

Transmembrane; Glycoprotein;

Repeat;

Signal;

Polymorphism;

TNFR_NGFR_1; 1.
TNFR_NGFR_2; 1.

SMART; SMO PROSITE; P

00020; TNFR 100208; TNFR PS00652; TN

TNFR;

7165; P:signal transduction; IPR001368; TNFR_c6.

; TNFR_c6.

PS50050;

Pfam; PF

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use by non-profimodified and this entities requires or send an email (
                                                                                                                     Genew; HGNC:11908; TNFRSF11A.

MIM; 603499; -.

MIM; 174810; -.

MIM; 602080; -.

GO; GO:0004872; F:receptor activity; TAS.

GO; GO:0007267; P:cell-cell signaling; TAS.

GO; GO:0008284; P:positive regulation of cell proliferation;

GO; GO:0007165; P:signal transduction; TAS.

InterPro; IPR001368; TNFR_c6.
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HSSP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Mutations in TNFRSF11A, familial expansile osteo Nat. Genet. 24:45-48(200-!- FUNCTION: Receptor f
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INTERACTI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ng B.R., Josien R., Lee S.Y., Vologo
oi Y.;
he TRAF family of signal transducer
the TRANCE receptor.";
Biol. Chem. 273:28355-28359(1998).
                                                                                                                                                                                                                                                                                                                                                                                                     similarities involvement thave involvement to pelvis and skill SIMILARITY: Co
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DISEA
osteo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCE
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P25942: 10
                                                                                                                                                                                                                                                                                                   non-profit institutions as long and this statement is not removed. requires a license agreement (See
                                                                                                                                                                                                                                                                                                                                         S-PROT entry is copyright. It is produced through a collaboration the Swiss Institute of Bioinformatics and the EMBL outstation - yean Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20082806; PubMed=10615125;
.E., Ralston S.H., Marken J., Book R.G.H., van Hul W., Whyte M.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the essential signaling receptor for osteoclast tiation factor in osteoclastogenesis.";
Biophys. Res. Commun. 253:395-400(1998).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                J-mediated osteoclastogenesis. Involved in the regulation of actions between T-cells and dendritic cells.

III: Interacts with TRAF1, TRAF2, TRAF3, TRAF5 and TRAF6.

ILULAR LOCATION: Type I membrane protein (Potential).

IE SPECIFICITY: Ubiquitous expression with high levels in tal muscle, thymus, liver, colon, small intestine and adrenal
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                                                                                                                                                                                                                                                                                                                                                                                                                    and skull
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ansile osteolysis.";
24:45-48(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                    [MIM:602080]; also known as familial Paget disease of 2 is a bone remodelling disorder with clinical ies to FEO. Unlike FEO, however, affected individuals lyement of the axial skeleton with lesions in the smith diskull.
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Higashio K.;
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R., Lee S.Y., V
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Q28203;
Q1-NOV-1997 (Rel. 35, Created)
Q1-NOV-1997 (Rel. 35, Last sequence update)
Q1-OCT-2003 (Rel. 42, Last annotation update)
Q28203;
Q1-NOV-1997 (Rel. 35, Created)
Q1-NOV-1997 (Rel. 35, Last sequence update)
Q1-NOV-1997 (Rel. 35, Last sequence update)
Q1-NOV-1997 (Rel. 35, Created)
Q1-NOV-1997 (Re
                                                                                                                                                     SEQUENCE FROM N.A.

MEDLINE=97281252; PubMed=9135560;
Hirano A., Brown W.C., Estes D.M.;
"Cloning, expression and biological homologue: role in B-lymphocyte grown both statements of the second statement of the second state
                                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Bovidae; Bovinae; Bos. NCBI_TaxID=9913;
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SIMILARITY:
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Similarity 31.7%;
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L -> LLLCALL (in FEO)
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R EMBL; U57745; Anc.
R HSSP; P25942; 1CDF.
R InterPro; IPR008063; Fas_reu.
JR InterPro; IPR001368; TNFR_c6.
DR Pfam; PF00020; TNFR_c6; 4.
DR PRINTS; PR01680; FASRECEPTOR.
SMART; SM00208; TNFR_NGFR_1; 1.
PROSITE; PS00652; TNFR_NGFR_2; 1.
PROSITE; PS50050; TNFR_NGFR_2; 1.

Transmembrane; Glycoprotein; Repeat; Signal.
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19 POTENTIAL.
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                                           TNR5 MOUSE STANDARD; PRT; 289 AA.

P27512; Q99NE0; Q99NE1; Q99NE2; Q99NE3;

01-AUG-1992 (Rel. 23, Created)

01-OCT-1996 (Rel. 34, Last sequence update)

10-OCT-2003 (Rel. 42, Last annotation update)

Tumor necrosis factor receptor superfamily member 5 precursor (CD40L receptor) (B-cell surface antigen CD40) (BP50) (CDw40)

TNFRSF5 OR CD40.
Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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Sciurognathi;
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                          Vertebrata; Euteleostomi;
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                  Muridae;
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                  Murinae;
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This SWISS-PROI CALLE Obetween the Swiss Institute of the European Bioinformatics I use by non-profit institute modified and this statement institute a license a license a license a license a
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SEQUENCE FROM N.A. (ISOFORM I).
MEDLINE=92105763; PubMed=1370315;
Torres R.M., Clark E.A.;
"Differential increase of an altern species of murine CD40 upon B lymph J. Immunol. 148:620-626(1992).
                                                entities or send a
                                                                                                                                                                                                                                                                                                                        Ishida T.
Yamamoto
"TRAF5, a
 EMBL;
EMBL;
EMBL;
EMBL;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   alternative Proc. Natl. [5]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              [4]
SEQUENCE
V).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRAIN=BALB/c;
Torres R.M.;
Submitted (SEP-
[3]
                                                                                                                                                                                                                                          family protein, mediates CD40 signaling.";
Proc. Natl. Acad. Sci. U.S.A. 93:9437-9442(1996)
-!- FUNCTION: Receptor for TNFSF5/CD40L.
-!- SUBUNIT: Interacts with TRAF3 and TRAF5. Int
TRAF2 AND TRAF6 (By similarity).
-!- SUBCELLULAR LOCATION: Type I membrane proteinand V); secreted (isoform II).
-!- ALTERNATIVE PRODUCTS:
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Howard M.
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Tone M., Tone Y.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A. (ISOFORM 1).
STRAIN=BALB/c; TISSUE=Liver;
STRAIN=93094586; PubMed=1281194;
MEDLINE=93094586; PubMed=1281194;
                                                                                                                                                                                                                                                                                                                                                                 INTERACTION WITH TRAF5
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SIMILARITY:
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Name=III;
IsoId=P27512-3;
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M83312;
M94126;
M94129;
M94128;
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                                                                                  38-PROT entry is copyright. It is produced through a copyright the Swiss Institute of Bioinformatics and the EMBL bean Bioinformatics Institute. There are no restrict
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                                                                                                                                                                                                                                                                                                                                                                                                                                           117110; PubMed=11172023; one Y., Fairchild P.J., Wy}n of CD40 function by its ite splicing.";
Acad. Sci. U.S.A. 98:1751
AAB08705.1;
AAA37404.1;
AAA37404.1;
AAA37404.1;
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Contains 4 TNFR-Cys repea
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Aoki T., Kobayashi
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PIR; A464
HSSP; P25
MGD; MGI:
InterPro;
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SIGNAL
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Similarity 31.0%;
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PS50050;
               KDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTC
                                RHCEPNOGLRVKKEGTAESDTVCTCKEGQHCT--
                                                RPCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPG-
                                                               TCSD--KQYLHDGQ--CCDLCQPGSRLTSHCTALEKTQCHPCDSGEFSAQWNREIRCHQH
                                                                                TCRDQEKEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWNYLTICQLC
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IPR001368; TNFR_c6.
)020; TNFR_c6; 4.
R01680; FASRECEPTOR.
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37; CAC29427.1;
38; CAC29428.1;
39; CAC29429.1;
90; CAC29430.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TNFR_NGFR_1;
TNFR_NGFR_2;
CHPCPVGFFSNQSSLFEKCYPWTSCEDKNLEVLQKGTSQTNVIC
                                                                                                                                                   289
                                                                                                                                                                  222
                                                                                                                                                                                  289
                                                                                                                                                                                                          234
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215
289
60
103
1144
187
37
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77
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119
1116
1143
153
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                                                                                                                                                                                                                                                   289
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                                                                                                                                  32111 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JOINED.
                                                                                                22;
                                                                                                                                                                /FTId=VSP_006477.
Missing (in isoform I
/FTId=VSP_006478.
KKVVKKP -> SGQETKG (i
                                                                                                                                                                                                                                                                                        BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                         POTENTIAL.
TUMOR NECROSIS FACTOR
SUPERFAMILY MEMBER 5.
EXTRACELLULAR (POTENT:
POTENTIAL.
CYTOPLASMIC (POTENTIAL
TNFR-CYS 1.
                                                                                                                                                                                                                                                                                                                                                  TNFR-CYS 4.
BY SIMILARITY.
BY SIMILARITY.
                                                                                                        Score
Pred.
                                                                                                                                Missing (in isofor
/FTId=VSP_006480.
C791CB6D2FEA574E
                                                                                                                                                                                                                                                                                                                                                                          TNFR-CYS
                                                                                                                                                                                                                                 Missing (in isoform II).
/FTId=VSP_006475.
GLKSRMRALLVIPVVMGILITIFGVFLYIK
                                                                                                                                                                                               KKVVKKPKDNEMLPPAARR
(in isoform III).
                                                                                                                                                                                                                         isoform
                                                                                                                                                                                                                                                                         N-LINKED (GLCNAC. . .) (POTENTIAL).
SCEDKNLEVLQKGTSQTNVICGLKSRMRALLVIPVVMG
RFKVPDASPAGHSCRDGHPHHHFRGVSLYQKGGQETKG
                                                                                                                                                        FTId=VSP
                                                                                                                                                                                                                                                         (in isoform
                                                                                                                                                                                                                  FTId=VSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.
                                                                                                ore 243.5; DB 1
ed. No. 5.6e-14;
Mismatches 81
                                                                                                                                                                                                         V).
SP_006476.
                                                                                                                                      (in isoform SP_006480.
                                                                                                                                                         006479.
                                                                                                                                                                                                                                                          006474.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                   (POTENTIAL)
                                                                                                               DB 1;
                               SKDCEACAOHTPCIPGFGVMEMATET
                                                                                                                                                                                                                                                                                                                                                                                                                   (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                   FACTOR RECEPTOR
                                                                                                81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Signal;
                                                                                                                                                                (in
                                                                                                                                 CRC64;
                                                                                                                                                IV).
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                                                                                               17;
                                                                                                                                                                 IV)
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                                                -TEAEL
                                                                                               Gaps
                183
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is not removed.

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RESULT
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SEQUENCE FROM N.A., AND VARIANTS
AND ARG-296.
STRAIN=129/Ola, and NIH Swiss; TJ
MEDLINE=98382527; PubMed=9714833;
Mizuno A., Murakami A., Nakagawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Davy E., Bucay N., Renshaw-Campbell P., Sander S., Van Suggs S., Boyle W.J.; "Osteoprotegerin: a novel s of bone density."; Cell 89:309-319(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOUSE
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28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
28-FEB-2003 (Rel. 41, Last annotation update)
Common necrosis factor receptor superfamily member (Osteoprotegerin) (Osteoclastogenesis inhibitory for the composition of the composi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
STRAIN=BALB/c; TISSUE=Kidney;
MEDLINE=97262071; PubMed=9108485;
MEDLINE=97262071; PubMed=9108485;
Simonet W.S., Lacey D.L., Dunstan C.R., Kelley M., Chang Simonet W.S., Lacey D.L., Boone T. Luethy R., Nguyen H.Q., Wooden S., Bennett L., Boone T. Luethy R., Nguyen H.Q., Wooden S., Tan H.-L., Trail G Derose M., Elliott R., Colombero A., Tan H.-L., Trail G Davy E., Bucay N., Renshaw-Gegg L., Hughes T.M., Hill D Davy E., Bucay N., Renshaw-Gegg L., Tarpley J., Derby P., L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
NCBI_TaxID=10090;
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SEQUENCE FROM N.A.
STRAIN=BALB/c; TIS
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Van G.,
Simonet
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Gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             resembling osteoclastogenesis."
J. Exp. Med. 192:463-474(2000).
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"Osteoprotegerin reverses
between the
the European
use by non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteoclasts and prevents vascul
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mizuno A., M
Higashio K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FUNCTION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Structure
                                                                          This
                                                                                                       whereas expression decreases at day 11 and increases from to 17. On day 15 found in developing bone primordia, brachiocephalic artery and ductus arteriosus, left main abdominal aorta and midgut.

INDUCTION: Upregulated by TGF-beta and estrogens. Downrown of the primordia of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FUNCTION: Acts as decoy receptor for RANKL and thereby neutralizes its function in osteoclastogenesis. Inhibits the activation of osteoclasts and promotes osteoclast apoptosis in vitro. Bone homeostasis seems to depend on the local RANKL/OPG ratio. May also play a role in preventing arterial calcification. May act as decoy receptor for TRAIL and protect against apoptosis. TRAIL binding blocks the inhibition of osteoclastogenesis.
                                                                                                                                                                                                                                                                                                                                                        SUBUNIT: Homodimer.
SUBCELLULAR LOCATION: Secreted.
SUBCELLULAR LOCATION: Highly expressed in liver, lung,
TISSUE SPECIFICITY: Highly expressed in decidua
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOUSE
                                                                                                                                                                                                                                                                                                                          and in embryo.
DEVELOPMENTAL STAGE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cture of the mouse and its expression 215:339-343(1998).
                                                                            SWISS-PROT
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Kaufman S.,
W.S.;
                            Bioinformatics
                                                  Swiss
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                                                  entry is copy
ss Institute
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, Kostenuik
  institutions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      osteoclastogenesis in embryogenesis."
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                                                  copyright.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteoporosis by inhibiting endostea rascular calcification by blocking a
                             e of Bioinformatics
Institute. There
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P.J., La
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Craniata; Vert Sciurognathi;
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hi; Muridae;
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                                                         produced through a collaboration atics and the EMBL outstation -
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        no restricti content is
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Murinae; Mus
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Shimamoto
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              modified entities
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PROSITE;
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REPEAT
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                /PCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCKNPLE
                                  )SVKQ--ECNRTHNRVCECEEGRY-
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CKCPDGF
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larity 29.6%;
Conservative 2
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:extracellular
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DEATH DOMAIN;

TNFR NGFR 1; 1

TNFR NGFR 2; 2
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FSGETSSKAPCIKHTNCSTFGLLLIQKGNATHDNVCSGNRE
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                                                                                                                                                                                                                                                                                                                                                                                                                     Glycoprotein; Repeat; Signal; Polymorph
21
BY SIMILARITY.
01
TUMOR NECROSIS FACTOR RECEPTOR
SUPERFAMILY MEMBER 11B.
62
TNFR-CYS 1.
65
TNFR-CYS 2.
                                                                                                                                                 45923 MW;
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JOINED.
JOINED.
JOINED.
                                                                                                            Score 242.5; DB 1;
Pred. No. 9.4e-14;
25; Mismatches 83;
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TNFR-CYS 4.
DEATH 1.
DEATH 1.
DEATH 2.
INVOLVED IN DIMERIZATIVOLVED IN DIMERIZATIVOLVED IN DIMERIZATIVOLVED IN DIMERIZATIVOLVED IN LARITYOUS SUIMILARITYOUS SWISS).

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                                     -LEIEFCLKHRSCPPGS
                                                                                                                                                                                                                                                                                                                                                                           DIMERIZATION
                                                                                                                                                   CRC64;
                                                                                                                                                                                       129/OLA
                                                                                                                                                                                                                           129/OLA AND
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                                                                                                               Indels
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EN SEQUENCE FROM N.A., .....

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CTISSUE=Eye;

XX MEDLINE=22388257; PubMed=12477932;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

RA Klausner R.D., Collins F.S., Wagner L., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.

RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.

Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.

Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez J.

RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez J.

RA Butterfield Y.S. N., Touchman J.W., Green E.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,

RA Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;

TGeneration and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.",

Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                           ACCOCK REPORT RAPARA RAPARA RECENTARIA RECUTARIA RECENTARIA RECENTARIA RECENTARIA RECENTARIA RECENTARIA RECUTARIA RECENTARIA RECENTARIA RECENTARIA RECENTARIA RECENTARIA RECUTARIA RECENTARIA RECENTAR
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DT 10-OCT-2003 (Rel. 41, Last sequence update)
DT 10-OCT-2003 (Rel. 42, Last annotation update)
DE Tumor necrosis factor receptor superfamily member 11I
DE (Osteoprotegerin) (Osteoclastogenesis inhibitory fact
GN TNFRSF11B OR OPG OR OCIF.

OC Homo sapiens (Human).

C Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; En
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; I
NCBI_TaxID=9606;
NCBI_TaxID=9606;
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A Simonet W.S., Lacey D.L., Dunstan C.R., Kelley M., Chang M.-S. A Luethy R., Nguyen H.Q., Wooden S., Bennett L., Boone T., Shima Derose M., Elliott R., Colombero A., Tan H.-L., Trail G., Sull Davy E., Bucay N., Renshaw-Gegg L., Hughes T.M., Hill D., Patt Campbell P., Sander S., Van G., Tarpley J., Derby P., Lee R., Suggs S., Boyle W.J.;
T "Osteoprotegerin: a novel secreted protein involved in the reg of bone density.";
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TISSUE=Lung cancer;
MEDLINE=98151033; PubMed=9492069;
Yasuda H., Shima N., Nakagawa N., Mochizuki S.-I., Yano K., Fu Sato Y., Goto M., Yamaguchi K., Kuriyama M., Kanno T., Murakam Tsuda E., Morinaga T., Higashio K.;
T "Identity of osteoclastogenesis inhibitory factor (OCIF) and Tosteoprotegerin (OPG): a mechanism by which OPG/OCIF inhibits Endocrinology 139:1329-1337(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Morinaga T., Nakagawa N., Yasuda H., "Cloning and characterization of the osteoprotegerin/osteoclastogenesis-ir Eur. J. Biochem. 254:685-691(1998).
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the gene encoding human
s-inhibitory factor.";
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Shenmen C.M., Schuler G.D.,
, Schaefer C.F., Bhat N.K.,

Wang J., Hsieh F.,
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Hofbauer L.C., Neubauer A., Heufelder A.E.;
"Receptor activator of nuclear factor-kappaB ligand and osteoprotegerin: potential implications for the pathogenesis and treatment of malignant bone diseases.";
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TISSUE=P1
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J. Biol. (
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-!- FUNCTION: Acts as d
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T. Biol. Chem. 273:5117-5123(1998).
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Emery J.
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MEDLINE=98148058; PubMed=9478964;
Yamaguchi K., Kinosaki M., Goto M., Kobayashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dul
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erization of structural domains of human osteoclastogenesis
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S-PROT entry is copyright. It is produced through a collaboration the Swiss Institute of Bioinformatics and the EMBL outstation - ean Bioinformatics Institute. There are no restrictions on its non-profit institutions as long as its content is in no way and this statement is not removed. Usage by and for commercial requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          YION: Acts as decoy receptor for RANKL and thereby neutralizes function in osteoclastogenesis. Inhibits the activation of sclasts and promotes osteoclast apoptosis in vitro. Bone stasis seems to depend on the local RANKL/OPG ratio. May also a role in preventing arterial calcification. May act as decoy tor for TRAIL and protect against apoptosis. TRAIL binding is the inhibition of osteoclastogenesis.
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Biophys. Res. Commun. 234:137-142(1997).
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and expression of osteoprotegerin from Homo sapiens.";
nim. Biophys. Sin. 31:680-684(1999).
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EMBL; AB002146; BAA25910.1; -.
EMBL; AB0008821; BAA32076.1; -.
EMBL; AB0008821; BAA32076.1; JOINED.
EMBL; AC030155; AAH30155.1; -.
EMBL; AF134187; AAF20168.1; -.
HSSP; P25942; 1CDF.

Genew; HGNC:11909; TNFRSF11B.

MIM; 602643; -.
GO; GO:0005576; C:extracellular; TAS.
GO; GO:0005125; F:cytokine activity; TAS.
GO; GO:0004872; F:receptor activity; TAS.
GO; GO:0007165; P:signal transduction; TAS.
GO; GO:0001501; P:skeletal development; TAS.
InterPro; IPR000488; Death.
InterPro; IPR001368; TNFR_C6.
Pfam; PF00020; TNFR_C6; 3.
SMART; SM00005; DEATH; 1.
SMART; SM00208; TNFR; 4.
PROSITE; PS50017; DEATH DOMAIN; FALSE_NEG.
PROSITE; PS00652; TNFR_NGFR_1; 1.
PROSITE; PS00652; TNFR_NGFR_2; 2.
Recentor: Anontocic: Cluronrotein. Dencat. Ci
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Best Local
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Similarity 30.9%;
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PS00652; TNFR_NGFR_1; 1
PS50050; TNFR_NGFR_2; 2
Apoptosis; Glycoprotei
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PDGFFSNETSSKAPCRKHTNCSVFGLLLTQKGNATHDNI
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BY SIMILARITY.
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N-LINKED
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SUPERFAMILY
TNFR-CYS 1.
TNFR-CYS 2.
TNFR-CYS 3.
TNFR-CYS 4.
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/FTId=VAR_013439.
C->S: ABOLISHES DIMERIZATION.
MISSING: ABOLISHES DIMERIZATION
D -> A (IN REF. 1).
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es 84;
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RESULT

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SEQUENCE FROM TISSUE=Embryonic ir
TISSUE=97262071;
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28-FEB-2003
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Mammalia; Eut
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Luethy R., N
Derose M., E
Davy E., Buc
Campbell P.,
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Cell 89:309-3
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SMART; SM00005; DEA
SMART; SM00208; TNF
PROSITE; PS50017; D
PROSITE; PS00652; T
PROSITE; PS50050; T
CYtokine; Apoptosis
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                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suggs S., Boyle W
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HSSP;
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                                                                                                                                                                                                                                                                                           InterPro;
                                                                                                                                                                                                                                                                                                       erPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: Homodimer (By similarity).
SUBCELLULAR LOCATION: Secreted (By similarity).
INDUCTION: Upregulated by osteopontin.
SIMILARITY: Contains 4 TNFR-Cys repeats.
SIMILARITY: Contains 2 death domains.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               seems to depend on the local RANKL/OPG ratio. May also in preventing arterial calcification. May act as decoy for TRAIL and protect against apoptosis. TRAIL binding inhibition of osteoclastogenesis (By similarity) subunit: Homodimer (By similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION
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Jutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyonic intestine;
262071; PubMed=9108485;
3., Lacey D.L., Dunstan C.R., Kelley M., Chang M.-S.,
3., Lacey D.L., Dunstan C.R., Kelley M., Chang M.-S.,
3., Lacey D.L., Dunstan C.R., Kelley M., Chang M.-S.,
Nguyen H.Q., Wooden S., Bennett L., Boone T., Shimamoto G.,
Nguyen H.Q., Wooden S., Bennett L., Trail G., Sullivan J.,
Elliott R., Colombero A., Tan H.-L., Trail G., Sullivan J.,
1., Sander S., Van G., Tarpley J., Derby P., Lee R.,
1., Sander S., Van G., Tarpley J., Derby P., Lee R.,
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0005; DEATH; 1.
0208; TNFR; 4.
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(Rel. 41, Last seque)
(Rel. 41, Last annot)
sis factor receptor of
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12; 1CDF.
1PR000488;
1PR001368;
1PR001368;
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-319(1997).
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TNFR_NGFR_1; 1.
TNFR_NGFR_2; 2.
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, Last annotation update)
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DEATH 1.
DEATH 2.
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Sciurognathi;
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TUMOR NECROSIS FACTOR RECEPTOR
SUPERFAMILY MEMBER 11B.
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Best Loc
Matches
                                                             Zhang W., Wan T., Cao A.;
Submitted (MAY-1999) to the EMBL/Gen
[4]
SEQUENCE FROM N.A., AND VARIANTS ARG
MEDLINE=21629477; PubMed=11756979;
Struyf F., Posavad C.M., Keyaerts E.
Spear P.G.;
"Search for polymorphisms i
Nectin-1, and Nectin-2 in i
J. Infect. Dis. 185:36-44(2
                                                                                                                                                                                                                                                                                                         SEQUENCE FACT.

SEQUENCE FACT.

MEDLINE=97306336; Pubric

Tan K.B., Ni
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Eukaryota; Metazoa; (
Mammalia; Eutheria; )
NCBI_TaxID=9606;
[1]
SEQUENCE FROM N.A.
TISSUE=Cervical adence
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Q92956; Q8WXR1;
16-OCT-2001 (Rel
16-OCT-2003 (Rel
10-OCT-2003 (Rel
                                                                                                                                                                                                       Wang S., Gentz R., Yu G.-L., Harrop J., Lyn Porter T.G., Truneh A., Young P.R.;
"A newly identified member of the tumor neconsuperfamily with a wide tissue distribution lymphocyte activation.";
J. Biol. Chem. 272:14272-14276(1997).
                                                                                                                                                                                                                                                                                                                                                                                            TISSUE=Cervical adenocarcinoma; MEDLINE=97053782; PubMed=8898196; Montgomery R.I., Warner M.S., Lum I "Herpes simplex virus-1 entry into the TNF/NGF receptor family."; Cell 87:427-436(1996).
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16-OCT-2001 (Rel. 40, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Tumor necrosis factor receptor superfamily member
(Herpesvirus entry mediator A) (Tumor necrosis fac
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TNFRSF14 OR HVEM OR HVEA.
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; Q96J31; Q9UM65;
40, Created)
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3., Ni J., Oh K.-O.,
Yu G.-L., Harrop J
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Primates;
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Brownstein M.J., Usdin T.B., Toshiyuki S., Carminci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Radriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
Bukesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Balakesley R.W., Touchman J.W., Green E.D., Dickson M.C.,
Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
"Generation and initial analysis of more than 15,000 full-length fuman and mouse cDNA sequences.";

I Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903 (2002).
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INTERACTION WITH TRAF2 AND TRAF5.
MEDLINE=97298041; PubMed=9153189;
MEDLINE=97298041; PubMed=9153189;
Hsu H., Solovyev I., Colombero A., Elliott R., Kelley M., Boy Hsu H., Solovyev I., Colombero A., Elliott R., Kelley M., Boy "ATAR, a novel tumor necrosis factor receptor family member, "ATAR, a novel tumor necrosis factor receptor family member, through TRAF2 and TRAF5.";
through TRAF2 and TRAF5.";
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                                                                                                                                                                                                                                                                                   entry of several wildtype HSV strains of cells, and mediated HSV entry into activ SUBUNIT: Interacts with TRAF2, TRAF3 and SUBCELLULAR LOCATION: Type I membrane pr TISSUE SPECIFICITY: WIDELY EXPRESSED, WI IN LUNG, SPLEEN, AND THYMUS.
SIMILARITY: Contains 3 TNFR-Cys repeats.
                                                                                                                                                                                                                                                                                                                    TNFSF1/lymphotoxin-alpha. Involved in lymphocyte activation. InfSF1/lymphotoxin-alpha. Involved in lymphocyte activation. an important role in HSV pathogenesis because it enhanced the entry of several wildtype HSV strains of both serotypes into cells, and mediated HSV entry into activated human T cells. SUBUNIT: Interacts with TRAF2, TRAF3 and TRAF5.
SUBCELLULAR LOCATION: Type I membrane protein (Probable). TISSUE SPECIFICITY: WIDELY EXPRESSED, WITH THE HIGHEST EXPRESSED.
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          153978; AAF75588.1;
373877; AAL47717.1;
373878; AAL47718.1;
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1403268; PubMed=11511370;
Willis S.H., Whitbeck J.C., Krummenacher C.,
R.J., Wiley D.C.;
implex virus glycoprotein D bound to the human
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rus entry mediator, a member of the tumor necessity mediator, a member of the
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7306297; PubMed=9162022;
S.A., Ayres T.M., Skubatch M., Gray C.L., Rothe M.,
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AAD00505.1;
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GO; GO:0005027; F:NGF/TNF (6 C-domain) rec
GO; GO:0005027; F:NGF/TNF (6 C-domain) rec
GO; GO:0006955; P:cell surface receptor li
GO; GO:0006955; P:immune response; TAS.

InterPro; IPR008063; Fas receptor.

InterPro; IPR001368; TNFR c6.

PFAINTS; PR01680; FASRECEPTOR.

SMART; SM00208; TNFR, 3.

PROSITE; PS00652; TNFR NGFR 1; 1.

PROSITE; PS50050; TNFR NGFR 2; 2.

Receptor; Transmembrane; Glycoprotein; Ref
3D-structure.

SIGNAL 1 38 POTENTIAL.

SIGNAL 1 38 TUMOR NECROSI
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                                                                         Score 230.5;
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A Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
A Altschul S.F., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
A Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
A Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
A Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
A Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
A Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
A Raha S.S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
A Rodriguez A.C., Grimwood J., Schmutz J., Wyers R.M.,
A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M.,
A Butterfield Y.S.N., Krzywinski M.I., Skalska U., Smailus D.E.,
A Schnerch A., Schein J.E., Jones S.J.M., Marra M.A.;
Thuman and mouse cDNA sequences.";
Thuman and mouse cDNA sequences.";
Thuman and mouse cDNA sequences.";
-I.- FUNCTION: May activate NF-kappa-B and JNK and promote apoptosis.
-I.- FUNCTION: May activate with TRADD
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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MEDLINE=98378343; PubMed=9714541;
Pan G., Bauer J.H., Haridas V., W
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SEQUENCE FROM N.A.
TISSUE=Brain, and Colon;
MEDLINE=22388257; PubMed=124779;
MEDLINE=22388257; Feingold E.A.,
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                                                                          FUNCTION: May activate NF-kappa-B and JNK and promote apoptosis.
SUBUNIT: Associates with TRADD.
SUBCELLULAR LOCATION: Type I membrane protein (Probable).
TISSUE SPECIFICITY: Highly expressed in heart, brain, placenta, pancreas, lymph node, thymus and prostate. Detected at lower pancreas, lymph node, thymus and prostate. Detected at lower intestine, colon, spleen, bone marrow and fetal liver. Very low intestine, colon, spleen, bone marrow and fetal liver. Very low levels were found in adult liver and peripheral blood leukocytes.
SIMILARITY: Contains 4 TNFR-Cys repeats.
SIMILARITY: Contains 1 death domain.
CAUTION: It is uncertain whether Met-1 or Met-25 is the initiator.
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-PROT entry is copyright. It is produced through a collaboration
ne Swiss Institute of Bioinformatics and the EMBL outstation -
nn Bioinformatics Institute. There are no restrictions on its
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is factor receptor superfamily member 21 precursor (TNFR-h receptor-6) (Death receptor 6).
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01-OCT-2003 (TrEMBLrel. 25, Last annotation update)
01-OCT-2003 factor receptor-2.
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TNFR-2.

Paralichthys olivaceus (Flounder).

Paralichthys olivaceus (Flounder).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;

Acanthomorpha; Acanthopterygii; Percomorpha; Pleuronectiformes;

Pleuronectoidei; Paralichthyidae; Paralichthys.

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Park C., Kurobe T., Hirono I., Aoki T.;

"Cloning and characterization of cDNAs for two distinct tfactor receptor superfamily genes from Japanese flounder
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GO:0004872; F:receptor activity; IEA.erPro; IPR001368; TNFR_c6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CD40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TaxID=9823
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                                                                                                                                                                                                                                                                                                                                     81
                                                                                                                                                                                                                                                                                                                                                                                                                                                62;
                                                                                                                                                                                                                                                                                                                                                                                                                                               26.1%;
Similarity 36.9%;
52; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           KCKSNKGLQYAQRCSSTTRTGCVCKPGMYC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALE-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DEVĢĶGŅN--HCVPCKAGHFQŅTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCKNPLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ea (Pig).
n; Metazoa;
Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Д.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Д
                                                                                                                                                                                                                                                                                                                                   MGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPG
                                                                                                                                                                                                                                                                                                                                                                       KENQYPTNSRCCNLCPPGQKLVNHCTEVTETECLPCSSSEFLATWNREKHCHQHKYCDPN
                                                                                                                                                                                                                                                                                                                                                                                                         KEYYEPQHRIÇCSRCPPGTYVSAKCSRIRDTVCATÇAENSYNEHWNYLTICQLCRPCDPV
                                                                                                                                                                                                                                                    GNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTC
                                                                                                                                                                                                                                                                                              LGLQVQREGTSKTDTTCVCSEGHHCTNSA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRELIMINARY;
                                                        (TrEMBLrel. (TrEMBLrel. (TrEMBLrel.
                                                                                                                                    PRELIMINARY;
                                                                                                                                                                                                                         CEPCPVGFFSNVSSASEKCQPWTSCESKGLVEQRAGTNKTDVVC
                      factor recus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chordata; Craniata; Ver
Cetartiodactyla; Suina;
       Chordata;
                                          receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21,
21,
24,
                                                           19,
19,
24,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      response; IEA transduction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Score; Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Last
Last
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EMBL/GenBank/DDBJ
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Last
                                                                                              Created)
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         Craniata;
                                                                                                                                     PRT;
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4.
20D446E44AF93DD2
                                          type
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                                           annotation
                                                                                sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    295.5;
No. 5.4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vertebrata;
ina; Suidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 3.4e-24;
                                           (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AA
                                                                                                                                                                                                                                                                                                      CESCTLHSLCFPGLGVKQMATEVSDTI--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -IMDFDNPYCAECRNYSQCRAGYGVSLP
       Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              activity;
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on update)
                                                                                                                                     AA
                                                             update)
on update)
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Sus.
          Euteleostomi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         278;
                                                                                                                                                                                                                                                                     183
                                                                                                                                                                                                                                                                                                                                            TEAELKDEVGK
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Best Loc
Matches
 Best
                 Query
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NCBI_TaxID=10
[1]
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through th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART; SM
PROSITE;
PROSITE;
Receptor.
NON_TER
NON_TER
                                                                                                                                                                                                                  "Expression Submitted (N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; AF420214; AAL16021.1; -.
GO; GO:0004872; F:receptor activity;
InterPro; IPR001368; TNFR_c6.
Pfam; PF00020; TNFR_c6; 4.
SMART; SM00208; TNFR; 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRAIN=Sprague-Dawle
Osburg B., Peiser C.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Q80WY6;
Q80WY6;
01-JUN-2003
01-JUN-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
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                                                    Pfam; PF0
SMART; SM
PROSITE;
PROSITE;
Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                       [1]
                                                                                                                                                                       GO; GO:000
InterPro;
                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=10116;
                                                                                                                                                                                                                                                                                                                                                                                                 Rattus norvegicus (Rat)
                                                                                                                                                                                                                                                                                                                                                                                                                     Tumor
                                                                                                                                                                                                           EMBL; AF498
                                                                                                                                                                                                                                                                                      STRAIN-Wistar;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Match
Local
                   Match
                                                                                                                                                                          GO:0004:
erPro; I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       g B., Peiser C., Doemling D., Schomburg L., Voigt K., receptors p60 and p80 are constitutively expressed by lary endothelial cells and participate in TNF-alpha trgh the blood-brain barrier.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               123
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                                                                                                                                                      PF000
                                                                                                                                                                                                                                                                                                                                                                                                                      necrosis factor receptor type II.
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                                                                                                                                    SMOO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          M00208; TN
PS00652;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PS00652; TNFR_NGFR_1;
PS50050; TNFR_NGFR_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PYASENQTCRDQEKEYYEPQHRICCSRCPPGTYVSAKCSRIRDTVCATCAENSYNEHWNY
   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PYKPEPGNQCQISQEYYDKKAQMCCAKCPPGQYAKHFCNKTSDTVCADCAAGMFTQVWNH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CKNPLEPLPPEMSGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PG
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                                                                                                                                                                                                                                                                   Α.,
                                                                                                                                                                                        A., Schafer M.K.;
on of TNFR2 in rat (
(MAR-2002) to the 1
)8039; AAP33151.1;
)4872; F:receptor a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               433
433 AA;
                                                                                             020; TNFR
0208; TNFR
500652; TN
550050; TN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SE-SPTPSAVPRT 195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        F-GVARSRTSNGNVICSACAPGTFSDTTSSTDVCRPHRICS--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ICQLC-RPC--DPVMGLEEIAPCTSKRKTQCRCQPGMFCA--AWALECTHCELLSDCP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25.7%; Score 291.5; DB 11; larity 33.3%; Pred. No. 2.3e-23; Conservative 24; Mismatches 91;
                                                                                                                                                                        PR001368; TNFR_c6.
                                                             474 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                        (TrEMBLrel. 24, Created)
(TrEMBLrel. 24, Last sequence update)
(TrEMBLrel. 25, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dawley;
                                                                                                                                                                                                                                                                                      TISSUE=Spleen;
                                                                                                                                     TNFR;
                                                                                                TNFR_NGFR_1;
TNFR_NGFR_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  45723 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            to the EMBL/GenBank/DDBJ
                                                                                                                                                        c6; 4.
                                                            50148 MW;
   25.7%;
                                                                                                                                        4
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                                                                                                                                                                                              activity;
                                                                                                                                                                                                                              dorsal root ganglion."
EMBL/GenBank/DDBJ data
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sciurognathi;
     Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                   Craniata; Vertebrata;
Sciurognathi; Muridae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75736D835E72CA4A CRC64;
                                                                                                  ωΝ
                                                              298C6AB9E8C8D714 CRC64;
     291.5; DI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IEA.
                                                                                                                                                                                                   IEA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        474
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                       Muridae;
         )B 11;
-23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   databases.
                                                                                                                                                                                                                                        databases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                                       Euteleostomi; Murinae; Ra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   --ILAIPGNASTDAV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     433;
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gene.";

Mamm. Genome

EMBL; X76401; CA...

R PIR; 148854; 148854.

R PIR; 148854; 1NCF.

JR HSSP; P19438; 1NCF.

DR GO; GO:0007166; P:cell surface rece

DR GO; GO:0008220; P:necrosis; IMP.

InterPro; IPR001368; TNFR_c6.

Pfam; PF00020; TNFR_c6; 4.

PROSITE; PS00652; TNFR_NGFR_1; 2

PROSITE; PS50050; TNFR_NGFR_2; 7

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CD40 and CD40 Ligand cDNA Sequences.";
ed (JUL-2003) to the EMBL/GenBank/DDBJ databases.
Y333789; AAP86653.1; ".
274 AA; 30284 MW; 9723789A07FAB6DB CRC64;
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L., Zeng Y.;
pts of CD40 isoform in peripheral mononuclear cells.
(JAN-2003) to the EMBL/GenBank/DDBJ databases.
25405; AAO43990.1; -.
                                                         FROM N.A.
                                                                                                                                                   03 (TrEMBLrel. 24, Created)
03 (TrEMBLrel. 24, Last sequence update)
03 (TrEMBLrel. 25, Last annotation update)
rosis factor receptor superfamily member 5
                                                                                                                                                                                                                                                                                                                      LKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTC
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                                                                                     D=9606;
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                                                                                              Metazoa; Chordata;
Eutheria; Primates;
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Carnivora; Fissipedia;
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Last annotation updat
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                                                                                                Craniata; Vertebrata; Catarrhini; Hominidae;
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                                                                                                                                                                                                                            223 AA
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.a; Canidae;
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GO; GO:0005743; C:mitochondrial inner men
GO; GO:0005488; F:binding; IEA.
GO; GO:0004888; F:transmembrane receptor
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0006955; P:apoptosis; IEA.
GO; GO:0006955; P:immune response; IEA.
GO; GO:0006810; P:transport; IEA.
InterPro; IPR006209; EGF_like.
InterPro; IPR008063; Fas_receptor.
InterPro; IPR001993; Mitoch_carrier.
InterPro; IPR001993; Mitoch_carrier.
PFINTS; PR01680; FASRECEPTOR.
SMART; SM00208; TNFR_C6; 4.
PROSITE; PS01186; EGF_2; 1.
PROSITE; PS01186; EGF_2; 1.
PROSITE; PS00215; MITOCH_CARRIER; 1.
PROSITE; PS00652; TNFR_NGFR_1; 1.
PROSITE; PS00652; TNFR_NGFR_1; 1.
                           MEDLINE=98414512; PubMed=9740674;
Hurle B., Segade F., Rodriguez R., Ramos S.S.
"The Mouse Tumor Necrosis Factor Receptor 2 Gand Characterization of the two Transcripts."
Genomics 52:79-98(1998).
EMBL; Y14619; CAA74969.1; JOINED.
EMBL; Y14621; CAA74969.1; JOINED.
EMBL; Y14622; CAA74969.1; JOINED.
EMBL; Y14623; CAA74969.1; JOINED.
EMBL; Y14623; CAA74969.1; JOINED.
EMBL; Y14679; CAA74969.1; JOINED.
HSSP; Q92956; 1JMA.
GO; GO:0004872; F:receptor activity; IEA.
InterPro; IPR001368; TNFR_C6.
Pfam; PF00020; TNFR_C6; 4.
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P80 TNF-alpha
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SEQUENCE
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            SMART; SI
PROSITE;
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                                                                                                                                                                                                                             Mus musculus (Mouse).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
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  SM00208; TN
TE; PS00652;
TE; PS50050;
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62; Conserv
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(TrEMBLrel.
ha receptor.
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TNFR_NGFR_1; 1.
TNFR_NGFR_2; 4.
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TNFR_NGFR_2;
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Sciurognathi;
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Mismatches
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annotation update)
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hi; Muridae;
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Best Local S
Matches 56
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GO; GO:0016020; C:membrane; IEA.
GO; GO:0004888; F:transmembrane receptor
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0006955; P:immune response; IEA.
GO; GO:0007165; P:signal transduction; IE
InterPro; IPR008063; Fas receptor.
InterPro; IPR001368; TNFR c6.
Pfam; PF00020; TNFR c6; 4.
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SEQUENCE
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"Cloning and
of the
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Q8WMQ2;
01-MAR-2002
01-MAR-2002
01-JUN-2003
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Bovidae; Ca
NCBI_TaxID=
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Ovis arie
                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.

Zawitkowski M.S., Russ G.R., Krishnan R.;

"Cloning and expression of the ovine CD40 molecule and the inhibition of the mixed lymphocyte reaction by the ovine CD40-EGFP fusion protein.";

Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.
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NON_TER
SEQUENCE
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SMART;
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                                                                                                                                                                                         PR01680; rau-
                                                                                                                                                                                             SM00208;
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                                                                                                         23.4%; Score 265.5; DB
Similarity 33.7%; Pred. No. 1e-20;
6; Conservative 21; Mismatches
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PS50050; TNFR_NGFR_2;
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 HCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTC
                                           NGLEEIAPCTSKRKTQCRCQPGMFCAAWALECTHCELLSDCPPGTEAELKDEVGKG--N
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                     .GLRIQSEGTLNTDTTCVCDEGQHCTSHT--CESCTPHSLCLPGFGVK--
                                                                                                                                                     277 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                               Craniata; Vertebrata; Euteleostomi; actyla; Ruminantia; Pecora; Bovoidea
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                                                                                                                                                      619F28BEA18A0D29 CRC64;
                                                                                                                                                                                                                                                                                   receptor
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                                                                                                                                                                                                                                                                                     activity;
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                                                                                                                               DB 6;
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                                                                                                                                  Length 277;
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       183
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                            QIATGVLD
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RESULT
Q805B0
ID Q8
AC Q8
DT 01
DT 01
DT 01
DT 11
OC TN
OC G6
OC A1
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Best Loc
Matches
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STRAIN=C57BL/6; TISSUE=Thymus;
Benencia F., Conejo-Garcia J.R., Courrege
"Light regulation in a murine model of ov
Submitted (MAR-2003) to the EMBL/GenBank/
EMBL; AY264405; AAO89081.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0007165; P:signal transduction; II
InterPro; IPR008063; Fas receptor.
InterPro; IPR008063; Fas receptor.
InterPro; IPR001368; TNFR c6.
Pfam; PF00020; TNFR c6; 4.
PRINTS; PR01680; FASRECEPTOR.
SMART; SM00208; TNFR, 4.
PROSITE; PS00652; TNFR NGFR 1; 1.
PROSITE; PS00652; TNFR NGFR 1; 1.
PROSITE; PS50050; TNFR NGFR 2; 3.
                            Tumor necrosis
TNFR-II.
Gallus gallus
                                                        Q805B0;
Q805B0;
01-JUN-2003
01-JUN-2003
01-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mus musculus (Mouse)
Eukaryota; Metazoa; (
Mammalia; Eutheria; I
NCBI_TaxID=10090;
 Eukaryota; Me
Archosauria;
Gallus.
                                                                                                                                                                                                                                                                                                                                                Receptor;
SIGNAL
SEQUENCE
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Q80WM9;
01-JUN-2003
01-JUN-2003
01-OCT-2003
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N-2003 (TrEMBLrel.:
T-2003 (TrEMBLrel.:
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UN-2003 (TrEMBLrel.
CT-2003 (TrEMBLrel.
r necrosis factor red
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Similarity 34.8%;
53; Conservative
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(TrEMBLrel. 24, Last sequence upd)
(TrEMBLrel. 25, Last annotation upsis factor receptor superfamily.
(Mouse).
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                                                                                                 PRELIMINARY;
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           Aves;
                             (Chicken)
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                                                                                                                                                                                     KRGTHDQDTVCADCLTGTF-SLGGTQEECLPWTNC-SAFQQEVRRGTNSTDTTCS
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          ; Chordata;
Neognathae;
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Rodentia;
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30171
                                             .. 24, Created)
.. 24, Last sequence upon 25, Last annotation under ceceptor-II.
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14; Mis
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he model of ovarian
EMBL/GenBank/DDBJ o
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C4A7EAD8EFC0521D
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          Craniata;
; Galliforn
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                                                                                                                                                                                                                                                                                                         Mismatches
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          ormes;
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                   Vertebrata;
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                                                                  update)
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-20;
86;
                                                        update)
         Phasianidae; Phasian
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Murinae; Mu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      precursor
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         Phasianinae;
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Best Loc
Matches
                                                                                             Eukaryota;
Mammalia; F
NCBI_TaxID=
                                              Strausberg
Submitted
EMBL; BC02
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GO; GO:000
GO; GO:000
GO; GO:000
GO; GO:000
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Sayed A., He
"Molecular"
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RESULT
Q8K2X6
ID Q8
AC Q8
AC Q8
DT 01
DT 01
DT 01
DT 01
RP S1
RP G0
DR G0
DR G0
DR G0
DR G0
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Q8K2X6;
Q8K2X6;
01-OCT-2002 (TrEMBLrel
01-OCT-2002 (TrEMBLrel
01-OCT-2003 (TrEMBLrel
01-OCT-2003 (TrEMBLrel
Similar to tumor necro
Mus musculus (Mouse).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfam; PF00020; TNFR_C6; 3
PRINTS; PR01680; FASRECEP
SMART; SM00208; TNFR; 4.
PROSITE; PS00652; TNFR_NG
PROSITE; PS50050; TNFR_NG
Receptor.
SEQUENCE 462 AA; 50141
  NCBI_Tax1
[1]
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=White leghorn H-B15; TISSUE=Spleen;
Sayed A., Horuchi H., Furusawa S., Matsuda H.;
"Molecular cloning and characterization of chicken Tumor necfactor receptor-II (TNFR-II) and Tumor necrosis factor receptassociated factor-5 (TRAF-5) genes.";
Submitted (JAN-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; AB101004; BAC55966.1; -.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004888; F:transmembrane receptor activity; IEA.
GO; GO:0004888; F:transmembrane receptor activity; IEA.
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0006955; P:immune response; IEA.
GO; GO:0007165; P:signal transduction; IEA.
InterPro; IPR008063; Fas receptor.
InterPro; IPR001368; TNFR c6.
 90:000
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C029254; AAH29254.1; -.
0016020; C:membrane; IEA.
0005634; C:nucleus; IEA.
0005840; C:ribosome; IEA.
0003677; F:DNA binding; IEA.
0003735; F:structural constit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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d (MAY-2002)
a (MAY-2002)
                                                                                                              FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  LTICQLCR-PCDFVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALE-CTHCELLSDCPPGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ite leghorn H-B15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PS50050; TNFR_NGFR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         D=9031;
                                                                                                                                                    Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                              tumor necrosis
                                                                                                                                                                                            (TrEMBLrel. 22, Created)
(TrEMBLrel. 22, Last sequence update)
(TrEMBLrel. 25, Last annotation updatumor necrosis factor receptor superfa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22.3%; Score 253; DB 13; larity 29.4%; Pred. No. 3.9e-19; Conservative 24; Mismatches 100;
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TNFR_NGFR_2;
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  constituent
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                                                                                                                                                                                                             annotation update)
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  of.
                                                                                                                                                                                              superfamily,
 ribosome;
                                                                                   databases
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   IEA.
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                                                                                                                                                       Mus.
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RESULT
Q9XSZ8
ID Q9
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GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0006955; P:immune response; IEA.
GO; GO:0006412; P:protein biosynthesis; I
GO; GO:0007165; P:signal transduction; II
InterPro; IPR008063; Fas_receptor.
InterPro; IPR001005; Myb_DNA_binding.
InterPro; IPR001865; Ribosomal_S2.
InterPro; IPR001368; TNFR_C6.
Pfam; PF00020; TNFR_C6; 4.
PROSITE; PR01680; FASRECEPTOR.
SMART; SM00208; TNFR; 4.
PROSITE; PS00037; MYB_1; 1.
PROSITE; PS000652; TNFR_NGFR_1; 1.
PROSITE; PS00652; TNFR_NGFR_1; 1.
PROSITE; PS00652; TNFR_NGFR_2; 4.
green monkey kidney cells with a herpes green fluorescence protein.";
Virology 258:365-374(1999).
EMBL; AF147720; AAD37381.1; -.
HSSP; Q92956; 1JMA.
GO; GO:0016020; C:membrane; IEA.
GO; GO:0004888; F:transmembrane receptor
GO; GO:0006915; P:apoptosis; IEA.
GO; GO:0007165; P:immune response; IEA.
GO; GO:0007165; P:signal transduction; II
InterPro; IPR008063; Fas receptor.
InterPro; IPR001368; TNFR_c6.
Pfam; PF00020; TNFR_c6; 3.
PRINTS; PR01680; FASRECEPTOR.
SMART; SM00208; TNFR; 3.
PROSITE; PS00652; TNFR_NGFR_2; 2.
                                                                                                                                                                                                                              [1]
SEQUENCE FROM N.A.
TISSUE=Kidney;
MEDLINE=99296730;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Receptor.
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                             Q9XSZ8;
Q9XSZ8;
01-NOV-1999
01-NOV-1999
01-JUN-2003
                                                                                                                                                                                                                                                                                    Cercopithecus aethiops (Green monkey) (Gr. Eukaryota; Metazoa; Chordata; Craniata; V. Mammalia; Eutheria; Primates; Catarrhini; Cercopithecinae; Cercopithecus.
                                                                                                                                                                                                                                                                                                                                        HVEAS
                                                                                                                                                                                                      MEDLINE=99296730; PubMed=10366573; Foster T.P., Chouljenko V.N., Kousoulas "Functional characterization of the Hve."
                                                                                                                                                                                                                                                                                                                                                    HveAs.
                                                                                                                                                                                                                                                                              Cercopithecinae;
NCBI_TaxID=9534;
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Similarity 31.0%;
54; Conservative
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Pred. No. 2.6e
22; Mismatches
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Last sequence up
Last annotation
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No. 2.6e-1
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.ni; Cercopithecidae;
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RESULT
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Submitted:
EMBL; X94355; CAD90722...
HSSP; Q92956; 1JMA.
GO; GO:0004872; F:receptor activity; II
GO; GO:0004872; F:receptor activity; II
InterPro; IPR001368; TNFR_c6.
Pfam; PF00020; TNFR_c6; 2.
SMART; SM00208; TNFR_ NGFR_1; 2.
PROSITE; PS00652; TNFR_NGFR_1; 2.
PROSITE; PS50050; TNFR_NGFR_2; 2.
PROSITE; PS50050; TNFR_NGFR_2; 2.
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01-AUG-1998
01-JUN-2003
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Shchelkunov
"Genes of a
                                                                                                                                                                                                                                                                                                                                SEQUENCE FRO
STRAIN=GRI-9
MEDLINE=9706
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Cowpox
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STRAIN=GRI-9
Shchelkunov
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SEQUENCE FRO
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                                                                                                                                                                                                   Ryazankina O.I., Petrov N.A., Gutorov V.V., Kotwal G.J., Sandakhchiev L.S.; "Structure-function and organization of cowpox virus strain GRI-90
                                                                                                                                                                              "Structure-function a complete genome."; Submitted (MAR-2003)
                                                                                                                                 Totmenin A.
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                                                                                                                                                                                                                                                                                    68532; PubMed=8963248;
F., Petrov N.A., Riazankina O.I., Totmen S.N., Sandakhchiev L.S.; circle of hosts for the cowpox virus."; Nauk 349:829-833(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21.4%; Score 242.5; DB 6; larity 32.2%; Pred. No. 3.3e-18; Conservative 18; Mismatches 85;
                                                                                                                                             90;
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2.
D2342F1040A00AE3
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                   CRC64;
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RESULT
Q712H3
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AC Q7
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DT 01
DT 01
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OC Pr
OC Pr
OC Pr
CO OC Pr
CO OC Pr
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OC Pr
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Search completed: August Job time : 119 secs
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Best Local S
Matches 48
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Best Local S
Matches 50
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

TISSUE=Head kidney;

Zou J., Secombes C.J.;

"Molecular cloning and expression analysis of a in rainbow trout, Oncorhynchus mykiss.";

Submitted (NOV-2002) to the EMBL/GenBank/DDBJ da EMBL; AJ517804; CAD57165.1; -.

Receptor.

SEQUENCE 318 AA; 35254 MW; 78F8135011283B43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q7T2H3 PRELIMINARY; PRT
Q7T2H3;
01-OCT-2003 (TrEMBLrel. 25, Creat
01-OCT-2003 (TrEMBLrel. 25, Last
01-OCT-2003 (TrEMBLrel. 25, Last
Tumour necrosis factor receptor.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleos:
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus
NCBI_TaxID=8022;
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Similarity 29.1%;
50; Conservative 2
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Pred. No. 4e-18;
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US-08-795-445A-48

US-08-795-446B-48

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US-08-795-446B-48

US-08-577-788C-48

US-09-580-235-2

US-09-580-181-2

US-09-580-181-8

US-09-580-181-8

US-09-580-181-8

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LENGTH:
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               MOLECULE
8-505-606-
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equence 4, Ap	equence 6. Appl	equence 4, Appl	equence 6, Appl	equence 4, Appl	equence 3, App	equence 1, Appl	e 4, Appl	equence 1, App	0. 539576	e 3, Appl	e 2, Appl	e 2, Appl	e 2, App	e 4, Appl	equence 2, Appl	e 4, Ap	Sequence 3, Appli

ALIGNMENTS

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Sequence 1, Applicaci
Patent No. 5925351
GENERAL INFORMATION:
                                                                                                                                                                          CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/505
FILING DATE: 21-JUL-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/378
FILING DATE: 26-JAN-1995
ATTORNEY/AGENT INFORMATION:
NAME: HALEY, Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B191
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9090
TELEFAX: (212) 596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. H.
STREET: 1251 Avenue of
CITY: New York
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                          R READABLE FORM:
R TYPE: Floppy disk
TTER: IBM PC compatible
TING SYSTEM: PC-DOS/MS-DOS
RARE: PatentIn Release #1.0, V
APPLICATION DATA:
                                                                                                                               X: (212) 596-9090
14-8367
NN FOR SEQ ID NO:
CHARACTERISTICS:
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1251 Avenue of the
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, James F.
FR: 27,794
FR: 8191
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N, Christopher D.
Paula S.
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Length 197;
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RESULT 3
US-09-303-262-1
; Sequence 1, Application US/09303262
; Patent No. 6669941
; GENERAL INFORMATION:
    APPLICANT: BROWNING, Jeffrey I BENJAMIN, Christoph
                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Soluble Lymphotoxin-B Receptors and Anti-lymphot TITLE OF INVENTION: Receptor and Ligand Antibodies, as Therapeutic A TITLE OF INVENTION: for the Treatment of Immunological Disease. FILE REFERENCE: B191
CURRENT APPLICATION NUMBER: US/09/000,166A
CURRENT FILING DATE: 1998-06-08
EARLIER APPLICATION NUMBER: PCT/US96/12010
EARLIER FILING DATE: 1996-07-19
NUMBER OF SEQ ID NOS: 3
SOFTWARE: Patentin Ver. 2.0
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          T: BROWNING, BENJAMIN, HOCHMAN, P. INVENTION: SO
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3, Jeffrey L.
N, Christopher D.
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: SOLUBLE LYMPHOTOXIN-BETA RE
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RESULT 4
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; Sequence 14, Application US/08820000
; Patent No. 6469144
; Patent No. 6469144
; GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi J.
TITLE OF INVENTION: Apo-2 L.
TITLE OF SEQUENCES: 28
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CORRESPONDENCE ADDRESS:
ADDRESSE: James F. Haley, Jr.
STREET: 1251 Avenue of the Americas
CITY: New York
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10020
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: PC-DOS/MS-DOS
SOFTWARE: PC-DOS/MS-DOS
SOFTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/09/303,262
FILING DATE: 30-Apr-1999
CLASSIFICATION NUMBER: US/08/505,606
FILING DATE: 21-JUL-1995
APPLICATION NUMBER: US 08/378,968
FILING DATE: 26-JAN-1995
AFTORNEY/AGENT INFORMATION:
NAME: HALEY, Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: B191
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 596-9000
TELEX: 14-8367
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LEWISCH! 197 amino acids
                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local
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TOPOLOGY: linear
TULE TYPE: peptide
NCE DESCRIPTION: SEQ ID
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E: amino acid
                                                                                                                                                                                                                                                                                                                                                                           100.0%; Score 1133; DB 4; larity 100.0%; Pred. No. 2.5e-99; Conservative 0; Mismatches 0;
                                                                                       US/08828683A
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DISEASE
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NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genented

Genentech,

Inc

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AND

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South

San

Francisco

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Way

California

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RESULT !
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                                                                                     APPLICANT: Ebner, Reinhard
APPLICANT: Yu, Guo-Liang
APPLICANT: Ruben, Steven M.
APPLICANT: Ullrich, Stephen
APPLICANT: Ullrich, Stephen
APPLICANT: Zhai, Yifan
TITLE OF INVENTION: Apoptosis Inducing Molec
FILE REFERENCE: 1488.065000C
CURRENT APPLICATION NUMBER: US/09/523,323
CURRENT FILING DATE: 2000-03-10
EARLIER APPLICATION NUMBER: 60/168,380
EARLIER FILING DATE: 1999-12-02
EARLIER FILING DATE: 1999-12-02
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Best Local S
Matches 170
                                                                                                                                                                                                                                                                                Sequence 57, Application US/09523323 Patent No. 6635743 GENERAL INFORMATION:
                                             EARLIER
EARLIER
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EARLIER
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08-828-683A-14
                                EARLIER
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APPLICATION NUMBER: US/08/828,683A

FILING DATE: 31-Mar-1997

CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/625328

FILING DATE: 1-Apr-1996

APPLICATION NUMBER: 08/710802

FILING DATE: 23-Sep-1996

ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P100
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
ORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE
MEDIUM TYPE:
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         APPLICATION NUMBER: 60/148,326 FILING DATE: 1999-08-11 APPLICATION NUMBER: 60/142,657 FILING DATE: 1999-07-06 APPLICATION NUMBER: 60/137,457
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CITY: Sou
STATE: Ca
COUNTRY:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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SOFTWARE: WinPatin
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RESULT
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Matches
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                                                                                                                                                                                                                                                                         LENGTH: 170
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: UNSURE
LOCATION: (7)
OTHER INFORMATION: May be any an
                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SOFTWARE:
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R FILING DATE: 1999-03-11
R APPLICATION NUMBER: 09/252,656
R FILING DATE: 1999-02-19
R APPLICATION NUMBER: 60/075,409
R FILING DATE: 1998-02-20
R APPLICATION NUMBER: 09/027,287
R APPLICATION NUMBER: 09/027,287
R APPLICATION NUMBER: 09/003,886
R FILING DATE: 1998-01-07
R APPLICATION NUMBER: 08/822,953
R FILING DATE: 1997-03-21
R APPLICATION NUMBER: 60/013,923
R FILING DATE: 1996-03-22
R APPLICATION NUMBER: 60/030,157
R APPLICATION NUMBER: 60/030,157
R FILING DATE: 1996-10-31
R OF SEQ ID NOS: 70
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Similarity 99.4%;
9; Conservative
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                                       KGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSDTTCK 170
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US-09-006-353A-6 Sequence Patent No GENERAL INFO SOFTWARE: Paten
CURRENT APPLICATIO
APPLICATION NUMB
FILING DATE:
CLASSIFICATION: COMPUTER REAMEDIUM TYFICOMPUTER:
OPERATING APPLICANT TITLE OF NUMBER OF SEQUENCE:

CORRESPONDENCE ADDRESS:

ADDRESSEE: HUMAN GENOME SCIENT

OATO KEY WEST AVENUE APPLICANT APPLICANT: STREET: CITY: STATE: COUNTRY: ZIP: 208 No. 6, Application US/09006353A
5. 6261801
INFORMATION: RE: PatentIn Release #1.0, APPLICATION DATA: TYPE: Floppy disk

R: IBM PC compatible

ING SYSTEM: PC-DOS/MS-DOS ROCKVILLE 20850 YEI, YING-FEI
YU, GUO-LIANG
GENTZ, REINER
RUBEN, STEVEN
INVENTION: TUMOR NECROSIS AGENT TION NUMBER: READABLE FORM: 3 Sn INFORMATION: DATA: US/09/006,353A SCIENCES, Versi FACTOR RECEPTOR 5 INC

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RESULT 7
US-09-573-986-6
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Best Local
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APPLICANT: Wei, Yi
APPLICANT: Ni, Jia
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                                                                                                                                                                                                SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (301) 309-8512
INFORMATION FOR SEQ ID NO: (SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                        FILE REFERENCE: 1488.1280004
CURRENT APPLICATION NUMBER: US
CURRENT FILING DATE: 2000-05
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                       APPLICANT: Ruben, TITLE OF INVENTION:
                                                                                                                                                          TYPE: PRT
ORGANISM: Homo
09-573-986-6
                                                                                                                                                                                                                                                                           APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: li
MOLECULE TYPE:
9-006-353A-6
                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: BROOKES, ANDER REGISTRATION NUMBER: REFERENCE/DOCKET NUM TELECOMMUNICATION INFO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               68.0%;
Local Similarity 70.7%;
nes 135; Conservative 1
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                                                                                                                   68.0%;
Similarity 70.7%;
5; Conservative 1
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                                       YNEHWNHLSTCQLCRPCDIVLGFEEVAPCTSDRKAECRCQPGMSCVYLDNECVHCEEERL
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TION INFORMATION:
(301) 309-8504
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Pred. No. 7.1e-6
L4; Mismatches
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                                                                                                                     Score 771; DB 4; Pred. No. 7.1e-65; Mismatches 40;
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RESULT 9
US-09-627-775-3
; Sequence 3, Application; Patent No. 6682739
; GENERAL INFORMATION:
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US-08-866-545-
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                                                                                                                                                                                            ; MOLECULE TYPE: US-08-866-545-3
                                                                                                                                                      Query Match
Best Local S
Matches 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL
                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-493-4935
TELEFAX: 650-493-5556
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                  INFORMATION
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COMPUTER READAB
MEDIUM TYPE:
COMPUTER: IB
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APPLICANT:
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SOFTWARE: FastSEC
CURRENT APPLICATION
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NAME: COruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 00
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...PE: amino acid
STRANDEDNESS: sin TOPOLOGY: 1:
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READABLE FORM:
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Similarity 100.0%;
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CATION DATA:
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1155 Avenue of the Amer
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Murali,
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ANALOGUES DESIGNED FROM BINDING SITES OF TUMOR NECROSIS FACTOR RECEPTOR SUPERFAMILY AND THEIR
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Pred. No. 5.1e-
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APPLICANT:

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Mark

Application 682739

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CURRENT APPLICATION NUMBER: US
CURRENT FILING DATE: 2000-07-
PRIOR APPLICATION NUMBER: 60/
PRIOR FILING DATE: 1999-07-28
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
    Query
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6015, Patent No. 6015, Patent No. 6015, Parent INFORMATION:
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Best I
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PC-DOS/MS-DOS

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Ve:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/974,022
                                                                                   NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 227 amino acids
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                           MOLECULE 7
8-974-022-4
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ATTORNEY/AGENT INFORMATION:
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ADDRESSEE: Amgen Inc
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                                      TOPOLOGY: 1
LECULE TYPE:
                                                            TYPE: amino
STRANDEDNESS:
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David L.
ne, Frank .
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APPLICATION NUMBER:
FILING DATE:
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STATE:
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                                                                                                                                       TYPE:
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                                       26.9%; Score 305; DB 3; I imilarity 35.0%; Pred. No. 2.8e-21; ; Conservative 24; Mismatches 78;
               HWNYLTICQLCR---PCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALE-CTHCELLS
                                                                                                                                                                                                   Winter, Robert B. CE/DOCKET NUMBER: FOR SEQ ID NO:
                                                                                                                                                                                                                                   AGENT INFORMATION:
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California
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Calzone, Frank J.
Chang, Ming-Shi
Chang, Ming-Shi
Chang, Ming-Shi
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ino acid
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4; Mismatches 78;
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RESULT 13
US-08-974-186-48
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TOPOLOGY: lin
MOLECULE TYPE:
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NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 5
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MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0

CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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T: One Amgen Center I
Thousand Oaks
: California
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INVENTION:
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Sequence

48,

Application US/08974186

Sequence 48 Patent No.

48,

GENERAL INFORMATION:
APPLICANT: Boyle,
APPLICANT: Lacey,

APPLICANT:
APPLICANT:
TITLE OF IN

Calzone, Frank J.
Chang, Ming-Shi
INVENTION: OSTEOPROTEGERIN

Boyle, will Lacey, David L. Calzone, Frank J.

NUMBER OF S CORRESPONDE

SEQUENCES: 5 ENCE ADDRESS:

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US-08-795-446B-
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GENERAL
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Matches
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NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US/08/974,18
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ZIP: 91320-1789
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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PRIOR APPLICATION DATE
APPLICATION NUMBER:
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8, Application US/08795446B
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ZENTION:
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llarity 35.0%; Pred. No. 2.8e-21;
Conservative 24; Mismatches 78;
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R: US/08/974,186
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STREET:
CITY: Th
STATE: C
COUNTRY:

Thousand Oaks California

1840 Dehavilland Drive

91320-1789

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APPLICANT: Boyle, William
APPLICANT: Lacey, David
APPLICANT: Calzone, Frank
APPLICANT: Calzone, Frank
APPLICANT: Chang, Ming-Shi
TITLE OF INVENTION: Osteoprotegerin
FILE REFERENCE: A-378CIP
CURRENT APPLICATION NUMBER: US/08/706,945D
CURRENT FILING DATE: 1996-09-03
PRIOR APPLICATION NUMBER: 08/577,788
PRIOR FILING DATE: 1995-12-22
NUMBER OF SEQ ID NOS: 145
SOFTWARE: PatentIn version 3.1
SEQ ID NO 134
LENGTH: 227
TYPE: PRT
ORGANISM: Homo sapiens
US-08-706-945D-134
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NAME: Winter, Robert B.
REFERENCE/DOCKET NUMBER:
INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0
FILING DATE:
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TOPOLOGY: lir
MOLECULE TYPE:
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SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/795,44
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS
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Similarity 35.0%;
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                                                                                                  DCPPG----TEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAP 174
                                                                                                                       HWNYLTICQLCR---PCDPVMGLEEIAPCTSKRKTQCRCQPGMFCAAWALE-CTHCELLS 119
                                       GNASRDAVCTSTSPTRSMAP 211
                                                            GTAQSDTTC--KNPLEPLPP 192
          28,
           2004,
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                                                                                 CKPCAPGTFSNTTSSTDICRPHQICN---
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1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US07_NEW_PUB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/US09B_PUBCOMB.pep:*

13: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

14: /cgn2_6/ptodata/2/pubpaa/US10A_PUBCOMB.pep:*

15: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

16: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

17: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

18: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*
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	1133	100.0	9 1	13	US-10-003-211-1	Sequence 1, Appli
2	1133	00.	Ü	ø	-907-372-1	Sequence 19, Appl
ω	1133	00.	S	9	-768-779A	Sequence 6. Appli
4	1133	٠	W	10	17-372-1	Sequence 19, Appl
ហ	1133	00.	£	12	-10-087-19	Sequence 942, App
σ	1133	00.	ω	14	-10-291-480-	Sequence 6, Appli
7	1133	00.	S	15	69-300-	Sequence 17, Appl
80	1133	00.	Ü	15	-10-262-445-1	Sequence 133, App
9	1129	•	Φ	9	-907-372-	Sequence 1, Appli
10	1129	•	Θ	10	-09-917-37	Sequence 1, Appli
11	987	•	7	13	12-793-	Sequence 14, Appl
12	970	85.6	172	15	-10-375-680-5	Sequence 57, Appl
13	780	•	Ы	9	-948-018-1	Sequence 19, Appl
14	771	•	0	12	87-192-	Sequence 939, App
75	771	•	⊢ ³	9	-826-212-	Sequence 6, Appli

ALIGNMENTS

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GENERAL INFORMATION:

APPLICANT: Biogen, Inc.

APPLICANT: Browning, et al.

TITLE OF INVENTION: Soluble Lymphotoxin Beta Receptor and TITLE OF INVENTION: Soluble Lymphotoxin Receptor and Ligand Antibodies as TITLE OF INVENTION: Anti-Lymphotoxin Receptor and Ligand Antibodies as TITLE OF INVENTION: Therapeutic Agents for the Treatment of Immunological TITLE OF INVENTION: Diseases

FILE REFERENCE: A013US

CURRENT APPLICATION NUMBER: US/10/003,211

CURRENT FILING DATE: 2007/10-31

PRIOR APPLICATION NUMBER: PCT/US97/19436

PRIOR APPLICATION NUMBER: PCT/US97/19436

PRIOR APPLICATION NUMBER: 60/029,060

PRIOR FILING DATE: 1996-10-25

NUMBER OF SEQ ID NOS: 1

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 1

LENGTH: 197

TYPE: PRT

ORGANISM: Homo Sapien

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US-10-003-211-
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Matches 197
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TITLE OF INVENTION: TNF RECEPTOR 2
FILE REFERENCE: PC-0050 US
CURRENT APPLICATION NUMBER: US/09/
CURRENT FILING DATE: 2001-07-27
NUMBER OF SEQ ID NOS: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 197
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ :
SOFTWARE: PERL
SEQ ID NO 19
LENGTH: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ
                                                                                                                                                                                                     Sequence 6, Application US Patent No. US20020127637A1 GENERAL INFORMATION:
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lequence 19, A
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ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
COUNTY:
ZIP: 20850
ZIP: 20850
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: SYSTEM: PC-DOS/MS-DOS
                                                                                     NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: HUMAN GENOME
STREET: 9410 KEY WEST A
CITY: ROCKVILLE
STATE: MD
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Pred. No. 1.7
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RESULT 4
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                     Query Ma
Best Loc
Matches
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                                                                                                                                                                       APPLICANT: Lal, Preeti G.
APPLICANT: Warren, Bridget A.
APPLICANT: Warren, Bridget A.
TITLE OF INVENTION: TNF RECEPTOR 2 RELATED
FILE REFERENCE: PC-0050 US
CURRENT APPLICATION NUMBER: US/09/917,372
CURRENT FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 20
                                                                                                                                                             NUMBER OF SEQ 1
SOFTWARE: PERL
                                                                                      TYPE: PRT
ORGANISM:
FEATURE:
NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                       LENGTH: 435
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APPLICATION NUMBER: 09/086,58

FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:

NAME: KENLEY K. HOOVER

REGISTRATION NUMBER: 40,302

REFERENCE/DOCKET NUMBER: PF36

TELECOMMUNICATION INFORMATION:

TELEPHONE: (301) 309-8439

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APPLICATION NUMBER: US/09/768,779A
FILING DATE: 25-Jan-2001
CLASSIFICATION: <Unknown>
                                  Similarity
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TULE TYPE: protei
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No. US20030068619A1
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                        Score 1133; DB 10; Pred. No. 1.7e-88; Mismatches 0;
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RESULT 6
US-10-291-480-6
; Sequence 6, Application US/10291480
; Publication No. US20030100069A1
; GENERAL INFORMATION:
; APPLICANT: Ni, Jian
; APPLICANT: Moore, Paul
; TITLE OF INVENTION: Human Tumor Necre
; FILE REFERENCE: PF368C1D1
; CURRENT APPLICATION NUMBER: US/10/29:
; CURRENT FILING DATE: 2002-11-12
; PRIOR APPLICATION NUMBER: 09/768,779
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Morris, David W.
APPLICANT: Engelhard, Eric K.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND
TITLE OF INVENTION: CANCER
FILE REFERENCE: 529452000122
CURRENT APPLICATION NUMBER: US/10/087,192
CURRENT FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 09/747,377
PRIOR APPLICATION NUMBER: US 09/747,377
PRIOR APPLICATION NUMBER: US 09/798,586
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 2059
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 942
LENGTH: 435
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US-10-087-192-942
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Publication No. US20
GENERAL INFORMATION:
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Best Local
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TYPE: PRT
ORGANISM: Homo:
-10-087-192-942
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                                                            Necrosis
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PRIOR APPLICATION NUMBER: 09/086,582
PRIOR FILING DATE: 1998-05-28
PRIOR APPLICATION NUMBER: 60/048,020
PRIOR FILING DATE: 1997-05-29
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin version 3.1
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Best Local S
Matches 197
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ORGANISM:
-10-369-300-
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APPLICANT: Hancock, Wayne
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: IMMUNE
TITLE OF INVENTION: DISORDERS USING COMBINATION
FILE REFERENCE: 7853-255
                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/369,300 CURRENT FILING DATE: 2003-02-19 PRIOR APPLICATION NUMBER: 60/358,463 PRIOR FILING DATE: 2002-02-19
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RESULT 8
US-10-262-445-13;
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SOFTWARE: CUL-
SEQ ID NO 133
SEQTH: 435
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FILE REFERENCE: 21402-462D
CURRENT APPLICATION NUMBER: US/10/262,
CURRENT FILING DATE: 2002-10-01
PRIOR APPLICATION NUMBER: 60/327,454
PRIOR FILING DATE: 2001-10-05
PRIOR APPLICATION NUMBER: 60/327,917
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APPLICATION NUMBER: 60/
FILING DATE: 2001-10-09
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APPLICATION NUMBER: 60/343,629
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APPLICATION NUMBER: 60
FILING DATE: 2001-10-1
                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/328,056 FILING DATE: 2001-10-09
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                                                                                                                                                                                       APPLICATION NUMBER:
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F SEQ ID NOS: 133
CuraSeqLi
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Similarity 100.
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Rieger, Danie
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Giot, Lo
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Mezes, Peter
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; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyt
US-09-907-372-1
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Patent No. US20020068242A1
GENERAL INFORMATION:
APPLICANT: Lal, Preeti G.
APPLICANT: Warren, Bridget A.
                                 SOFTWARE:
SEQ ID NO 1
                                                                                                                                         Sequence 1, Applicat publication No. US20 GENERAL INFORMATION:
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                                                                                                       APPLICANT: Lal APPLICANT: Wa TITLE OF INVEN
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TITLE OF INVENTION: TNF RECEPTOR 2
FILE REFERENCE: PC-0050 US
CURRENT APPLICATION NUMBER: US/09/9
CURRENT FILING DATE: 2001-07-27
NUMBER OF SEQ ID NOS: 20
                                                      APPLICANT: Warren, Bridget A.
TITLE OF INVENTION: TNF RECEPTOR 2 RELATED
FILE REFERENCE: PC-0050 US
CURRENT APPLICATION NUMBER: US/09/917,372
CURRENT FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 20
          LENGTH:
TYPE: PI
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  ORGANISM:
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ilarity 99.5%;
Conservative
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; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Inc:
US-09-917-372-1
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US-10-112-793-14
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GENERAL INFORMATION:
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Best Local
                        10-11
                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 170 amino acids
                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Ashkenazi, Av
TITLE OF INVENTION: Apo-
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentec
Match
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
SOFTWARE: WinPatin (Genented
CURRENT APPLICATION DATA:
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COMPUTER READABLE
MEDIUM TYPE:
                                  SEQUENCE
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96; Conservative
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CITY: South San Fr
STATE: California
COUNTRY: USA
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                                                                                                                                           NAME: Marschang, Diane L. REGISTRATION NUMBER: 35,600 REFERENCE/DOCKET NUMBER: P1
                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/10, FILING DATE: 28-Mar-2002 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CPPGTEAELKDEVGKGNNHCVPCKAGHFQNTSSPSARCQPHTRCENQGLVEAAPGTAQSD
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                                              TYPE: Amino Acid
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Vo. US20020192729A1
                                   DESCRIPTION:
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TYPE: PRT

ORGANISM: Homo sapiens

FEATURE:

NAME/KEY: misc_feature

LOCATION: (7)..(7)

OTHER INFORMATION: Xaa can be as

US-10-375-680-57
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US-10-375-680-
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US-09-948-018
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Matches 169
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Matches
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Patent No. US20020150977A1

GENERAL INFORMATION:
APPLICANT: Theill et al
TITLE OF INVENTION: TNF RECEPTOR-LIKE MOLECULES
FILE REFERENCE: 01017/37677
CURRENT APPLICATION NUMBER: US/09/948,018
CURRENT FILING DATE: 2001-09-05
PRIOR APPLICATION NUMBER: US 60/230,191
PRIOR FILING DATE: 2000-09-05
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TITLE OF IN
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CURRENT FILING DATE: 2003-02-28
PRIOR APPLICATION NUMBER: US 60/360,234
PRIOR FILING DATE: 2002-03-01
NUMBER OF SEQ ID NOS: 70
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                                                                                                         Application US/09948018
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Sequence 939, Application US/10087192; Publication No. US20020182586A1; GENERAL INFORMATION:
APPLICANT: Morris, David W.
APPLICANT: Engelhard, Eric K.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND ME
TITLE OF INVENTION: CANCER
FILE REFERENCE: 529452000122
CURRENT APPLICATION NUMBER: US/10/087,192
CURRENT FILING DATE: 2002-03-01
PRIOR APPLICATION NUMBER: US 09/747,377
PRIOR APPLICATION NUMBER: US 09/747,377
PRIOR APPLICATION NUMBER: US 09/798,586
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2001-03-02
NUMBER OF SEQ ID NOS: 2059
SOFTWARE: Fast SEO for Minds
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Best Local Similarity 70.7%;
Matches 135; Conservative 1
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SEQ ID NO 939
LENGTH: 402
TYPE: PRT
ORGANISM: Mus musculus
-10-087-192-939
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ORGANISM: Mus
-09-948-018-19
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                                                                                                                                                                                            Score 771; DB 12;
Pred. No. 1.2e-57;
14; Mismatches 40;
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RESULT 15 US-09-826-

Sequence

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Application

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; LENGTH: 415
; TYPE: PRT
; ORGANISM: Homo
US-09-826-212-6
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APPLICANT: Wei, Ying-Fei
APPLICANT: Gentz, Reiner
APPLICANT: Ruben, Steven
APPLICANT: Ni, Jian
ITILE OF INVENTION: Tumor Necrosis Factor Re
FILE REFERENCE: 1488.1280006
CURRENT APPLICATION NUMBER: US/09/826,212
CURRENT FILING DATE: 2001-04-05
NUMBER OF SEQ ID NOS: 26
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SDT
                        SDTTCKNPLEP 189
                                                                                              YNEHWNHLSTCQLCRPCDIVLGFEEVAPCTSDRKAECRCQPGMSCVYLDNECVHCEEERL
TICKNPPEP
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                                                                                                                                                                                             68.0%; Score 771; DB 9; larity 70.7%; Pred. No. 1.2e-57; Conservative 14; Mismatches 40
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Search completed: August 28, 2004, 01:53:06 Job time : 128 secs